

FIGURE

Leu Ala Ser Arg Cys Arg Cys Cys Arg Pro Leu Thr Leu Ser Phe Ala Leu Cys Trp Arg Val Glu Ala Ala Ala Ala Gly Arg Cys Arg ** His Phe His Trp Ala Gly Ala Cys Lys Pro Leu Pro Leu Val Glu Ala Ala Asp Thr Phe Ile Gly Leu 3. TGG TCG CGT GAA GCC GTC GCC GTC GTG GAG CCG TCG CAG TCA CTT TTA CGG TTC 9 18 27 36 45 54

S. ACC AGC GCA CTT CGG CAG CGG CAG CAC CTC GGC AGC GTC AGI GAA AAT GCC AAG

Thr Ser Ala Leu Arg Gln Arg Gln His Leu Gly Ser Val Ser Glu Asn Ala Lys Pro Ala His Phe Gly Ser Gly Ser Thr Ser Ala Ala Ser Val Lys Met Pro Ser Gln Arg Thr Ser Ala Ala Ala Ala Ala Pro Arg Gln Arg Gln 2. Lys Cys Gln Ala

Ser Phe Arg Gly Ala Val Gly Tyr Ser Thr Pro Thr Gly Tyr Asp Lys Leu Phe Ala Ala Arg Leu Gly Het Leu Pro Pro His Glu Gly Lys Ile Ile Arg Leu Phe Leu Pro Gly Cys Gly Trp Leu Leu His Thr Asn Val Arg Leu Leu Gly GTT CTT TTC GCC GGG CGT TGG GGT ATT CTC CAC CCA CAA GTG GGA ATT ATT AGG 63 72 81 99 108 CAA GAA AAG CGG CCC GCA ACC CCA TAA GAG GTG GGT GTT CAC CCT TAA TAA TCC Gln Glu Lys Arg Pro Ala Thr Pro *** Glu Val Gly Val His Pro *** Ser Lys Lys Ser Gly Pro Gln Pro His Lys Arg Trp Val Phe Thr Leu Asn Asn Pro Arg Lys Ala Ala Arg Asn Pro Ile Arg Gly Gly Cys Ser Pro Leu Ile Ile Leu

Arg Pro Pro Ser Phe Cys Phe Val Pro Ala Glu Leu Arg Gly Lys Gln Asn Asn Gly Leu Leu Leu Phe Val Phe Tyr Pro Leu Lys Trp Asp Gly Lys Lys Ile Ile Glu Ser Ser Ser Phe Phe Leu Ile Arg Ser Ser Gly Ile Glu Arg Lys Ser AAG GCT CCT CCT CTT TTT GTT TTA TGC CCT CGA AGG TTA GAG GGA AAA ACT AAT 117 126 135 144 153 162

TTC CGA GGA GGA GAA AAA CAA AAT ACG GGA GCT TCC AAT CTC CCT TTT TGA TTA Phe Arg Gly Gly Glu Lys Gln Asn Thr Gly Ala Ser Asn Leu Pro Phe Leu Ser Glu Glu Glu Lys Asn Lys Ile Arg Glu Leu Pro Ile Ser Leu Phe Asp Tyr Pro Arg Arg Arg Lys Thr Lys Tyr Sly Ser Phe Gln Ser Pro Phe Leu Ile Ile

Gln Lys His Arg Pro Leu Asn Pro Leu Pro Tyr Phe Glu Glu Gly Gly Pro Thr Lys Asn Thr Ala Leu Phe Thr Gln Phe Leu Thr Ser Ser Arg Vol Glu Leu Pro Lys Thr Gln Pro Ser Ser Pro Lys Ser Ser Pro Leu Val Gly *** Arg Trp Pro AAA ACA AAC ACC GCT CCT TCC AAA CCT TCT CCC ATC TTG AGG AGT GGA GGT CCC 171 180 189 198 207 216

TTT TGT TTG TGG CGA GGA AGG TTT GGA AGA GGG TAG AAC TCC TCA CCT CCA GGG Phe Cys Leu Trp Arg Gly Arg Phe Gly Arg Gly *** Asn Ser Ser Pro Pro Gly Phe Val Cys Gly Glu Glu Gly Leu Glu Gly Arg Thr Pro His Leu Gln Gly Leu Phe Val Ala Arg Lys Val Trp Lys Arg Val Glu Leu Leu Thr Ser Arg Gly

Gln Ser Asn Gln *** Ser Ala Ser Lys *** Cys Pro Ser Thr Thr Asn Gln His Lys Arg Ile Lys Ser Lyu Leu Leu Ser Lys Val Leu His Leu Pro Ile Lys Thr Asn Ala Phe Lys Ala Leu Phe Cys Val Lys Leu Leu Thr Phe His Tyr Lys Pro CAA ACG CTT AAA ACG ATT CTT CGT CTG AAA ATT GTT CCA CTT CAC CAT AAA ACC 225 234 243 252 261 270 GTT TGC GAA TTT TGG TAA GAA GCA GAC TTT TAA CAA GGT GAA GTG GTA TTT TGG Val Cys Glu Phe Cys *** Glu Ala Asp Phe *** Gln Gly Glu Val Val Phe Tro Phe Ala Asn Phe Ala Lys Lys Gln Thr Phe Asn Lys Val Lys Trp Tyr Phe Gly Leu Arg Ile Leu Leu Arg Ser Arg Leu Leu Thr Arg *** Ser Gly Ile Leu Val

Gly Ser Gly Cys Arg Ser Leu Ser Leu Phe Arg Gly Ala Ser Tyr Leu Ile Ser Gly Ala Ala Val Asp Leu Phe Arg Phe Ser Gly Val Leu Leu Ile Phe Phe Val Ala Arg Gln Trp Het Ser Phe Ala Phe Pro Val Ser Trp Cys Phe Leu Ser Tyr ACG GGC GAC GGT GTA GCT CTT TCG CTT TCC TTG GCT GGT CGT CTT ATT TCT TAT 279 288 297 306 315 324

TGC CCG CTG CCA CAT CGA GAA AGC GAA AGG AAC CGA CCA GCA GAA TAA AGA ATA Cys Pro Leu Pro His Arg Glu Ser Glu Arg Asn Arg Pro Ala Glu Arg Ile Ala Arg Cys His Ile Glu Lys Ala Lys Gly Thr Asp Gln Gln Asn Lys Glu Tyr Pro Ala Ala Thr Ser Arg Lys Arg Lys Glu Pro Thr Ser Arg Ile Lys Asn Thr

Cys Tyr Leu Leu Gly Cys Val --- Arg Thr His Leu Glu Ala Ser Gly Pro Ser Ala Thr Phe Phe Ala Val Tyr Lys Asp Leu Thr Ser Ser Arg Pro Val Leu Pro Gln Leu Leu Ser Pro Trp Het Ser Ile Ser His Pro Ala Gly Arg Phe Trp Pro GAC GTC ATT TCT TCC GGT GTA TGA ATA GCT CAC ACC TCG AGG CGC CTT GGT CCC 333 342 351 360 369 378

CTG CAG TAA AGA AGG CCA CAT ACT TAT CGA GTG TGG AGC TCC GCG GAA CCA GGG Leu Gln --- Arg Arg Pro His Thr Tyr Arg Val Trp Ser Ser Ala Glu Pro Gly Cys Ser Lys Glu Gly His Ile Leu Ile Glu Cys Gly Ala Pro Arg Asn Gln Gly Ala Val Lys Lys Ala Thr Tyr Leu Ser Ser Val Glu Leu Arg Gly Thr Arg Gly

Ala Cys Arg Gly Thr *** Gln Gln Ser Tyr Gly Lys Pro Ser Pro Thr Lys Pro Leu Ala Ala Val Gln Arg Ser Ser His Thr Gly Lys Gln Leu Arg Pro Arg Gln Phe Arg Leu Ser Arg Asp Val Ala Thr Leu Val Arg Lys Ser Val Pro Asp Lys CTT CGC GTC GCT GGA CAG ATG ACG ACA CTC ATG GGA AAA CCT CTG CCC CAG AAA 387 396 405 414 423 432 432 GAA GCG CAG CGA CCT GTC TAC TGC TGT GAG TAC CCT TTT GGA GAC GGG GTC TTT Glu Ala Gln Arg Pro Val Tyr Cys Cys Glu Tyr Pro Phe Gly Asp Gly Val Phe Lys Arg Ser Asp Leu Ser Thr Ala Val Ser Thr Leu Leu Glu Thr Gly Ser Leu Ser Ala Ala Thr Cys Leu Leu Leu Val Pro Phe Trp Arg Arg Gly Leu Trp

Ser Gln Leu Arg Ala Thr Glu Gln Leu Thr His Ser Phe Asn Gly Arg Ala Pro His Ser Tyr Gly Leu Leu Lys Arg Tyr Arg Ile His Ser Ile Glu Ala Pro Gln Thr Val Thr Ala Ser Cys Asn Gly Thr Val Tyr Thr Leu Phe Lys Arg Pro Ser CCA CTG ACA TCG GCT CGT CAA AGG ACA TTG CAT ACA CTC TTT AAA GGC GCC CGA 41 450 459 468 477 486 GGT GAC TGT AGC CGA GCA GTT TCC TGT AAC GTA TGT GAG AAA TTT CCG CGG GCT Gly Asp Cys Ser Arg Ala Val Ser Cys Asn Val Cys Glu Lys Phe Pro Arg Ala Val Thr Val Ala Glu Gln Phe Pro Val Thr Tyr Val Arg Asn Phe Arg Gly Leu Leu Pro Ser Ser Phe Leu Arg Met Glu Ile Ser Ala Gly Trp

Gln Val Lys Ser Leu Ser Arg Ser Ser Ala Ala Ala His Asn Ser Ser Leu Gln Ser Phe Lys Gln Phe His Ala Pro Leu His Leu Leu Thr Ile Pro Leu Cys Ser Ala Ser Ser Lys Phe Thr Leu Pro Phe Ile Cys Cys Arg Ser Gln Phe Val Ala CCG ACT TGA AAA CTT TCA CTC GCC CTT CTA CGT CGC ACT AAC CTT CTG TCG 495 504 513 522 531 540 GCC TGA ACT TTT GAA/AGT GAG CGG GAA GAT GCA GCA GCG TGA TTG GAA GAC AGC Gly *** Thr Phe Glu Ser Giu Arg Glu Asp Ala Ala Ala *** Leu Glu Asp Ser Ala Glu Leu Leu Lys Val Ser Gly Lys Met Gln Gln Arg Asp Trp Lys Thr Ala Leu Asn Phe *** Lys *** Ala Gly Arg Cys Ser Ser Val Ile Gly Arg Gln Leu

FIGURE 2 (continuation 1)

Val Arg *** Leu Pro Gly Ala Arg Asn His Ser Ser Gly Thr Pro Gly Tyr Asn Tyr Val Asp Tyr His Ala Arg Gly Thr Thr Pro Veu Ala Leu Pro Gly Thr Ile Thr Cys Thr Het Thr Pro Gly Gly Pro Gln Pro Phe Leu Trp His Ala Arg Leu ACA TGT GCA GTA TCA CCC GGG CGG GCC AAC ACC CTT CTC GGT CAC CCG GGC ATT S49 S58 S67 S85 S94 TGT ACA CGT CAT AGT GGG CCC GGC CGG TTG TGG GAA GAG CCA GTG GGC CCG TAA Cys Thr Arg His Ser Gly Pro Ala Arg Leu Trp Glu Glu Pro Val Gly Pro Val His Val Ile Val Gly Pro Pro Gly Cys Gly Lys Ser Gln Trp Ala Arg Asn Tyr Thr Ser *** Trp Ala Arg Pro Val Val Gly Arg Ala Ser Gly Pro Val Ile

Gln Gln Ala *** Pro Cys Arg Ser Ser Ala *** Tyr Phe Tyr Thr Thr Pro His Lys Ser Leu Arg Pro Val Gly Val Pro Leu Arg Thr Ser Ile Leu Pro Pro Ile Lys Ala Ser Gly Leu Ser Val *** Gln Phe Gly Leu Leu Phe Leu His His Ser AAA ACG ACT CGG ATC CCT GTG GAT GAC CTT CGG ATC ATC TTT ATT CAC CAC CCT 603 612 621 630 639 648

TTT TGC TGA GCC TAG GGA CAC CTA CTG GAA GCC TAG TAG AAA TAA GTG GTG GGA

Phe Cys *** Ala *** Gly His Leu Leu Glu Ala *** *** Lys *** Val Val Gly Phe Ala Glu Pro Arg Asp Thr Tyr Trp Lys Pro Ser Arg Asn Lys Trp Trp Asp Leu Leu Ser Leu Gly Thr Pro Thr Gly Ser Leu Val Glu Ile Ser Gly Gly Het

Ile Asp His Leu Leu Leu Gln Gln Lys Pro His Asn Lys His Ser Thr Val Lys Ser Ile Het Ser Phe Phe Asn Asn Asn Gln Ile Ile Lys Ile Ala Pro Arg Pro Tyr Pro Ser Ser Thr Thr Thr Lys Ser Ser Lys Pro Gln Asn Gly ACC TAT AGT ACC TCT TCT TCA ACA ACA AAA CCT ACT AAA AAT ACC GAC CAA TGG 657 666 675 684 693 702 TGG ATA TCA TGG AGA AGA AGT TGT TGT TTT GGA TGA TTT TTA TGG CTG GTT ACC Trp Ile Ser Trp Arg Arg Ser Cys Cys Phe Gly Phe Leu Trp Leu Val Thr Gly Tyr His Gly Glu Glu Val Val Leu Asp Asp Phe Tyr Gly Trp Leu Pro Asp Ile Het Glu Lys Lys Leu Leu Phe Trp Het Ile Phe Het Ala Gly Tyr Leu

Pro His Asp Val Ser Val Thr His Gly Thr Asp Het Ser Gln Leu Ser Leu Pro Ile Ile ** Gln Ser Gln Thr Val Pro Ile Trp Gln Ser Tyr Leu Ser Phe Gln Ser Ser Arg Ser Leu Ser His Ser Arg Tyr Gly Asn Val Thr. Ser Val Leu AAC CCT ACT AGA TGA CTC TGA CAC ACT GGC CAT AGG TAA CTG ACA TCT CTG ATT 711 720 729 738 747 756.

TTG GGA TGA TCT ACT GAG ACT GTG TGA CCG GTA TCC ATT GAC TGT AGA GAC TAA Leu Gly ** Ser Thr Glu Thr Val ** Pro Val Ser Ile Asp Cys Arg Asp ** Trp Asp Asp Leu Leu Arg Leu Cys Asp Arg Tyr Pro Leu Thr Val Glu Thr Lys Gly Het Ile Tyr ** Asp Cys Val Thr Gly Ile His ** Leu ** Arg Leu Lys

Pro Tyr Gln Glu Lys Lys Pro Gly Cys Tyr Lys Ser *** Trp Cys Asp Pro Gly Pro Thr Ser Asn Arg Lys Gln Gly Ala Thr Asn Gln Asn Gly Ala Ile Leu Gly Pro Pro Val Thr Gly Lys Lys Ala Arg Leu Ile Lys Ile Val Leu Leu *** Ala TCC CCC ATG ACA AGG AAA AAA CCG GGC GTC ATA AAA CTA ATG GTC GTT AGT CCG 765 774 783 792 801 810 AGG GGG TAC TGT TCC TTT TTT GGC CCG CAG TAT TTT GAT TAC CAG CAA TCA GGC Arg Gly Tyr Cys Ser Phe Phe Gly Pro Gln Tyr Phe Asp Tyr Gln Gln Ser Gly Gly Gly Thr Val Pro Phe Leu Ala Arg Ser Ile Leu Ile Thr Ser Asn Gln Ala Gly Val Leu Phe Leu Phe Trp Pro Ala Val Phe *** Leu Pro Ala Ile Arg Pro

FIGURE 2 (continuation 2)

Gly Pro Ile Thr Ser Arg Leu Gln Gln Gly Leu Gln Leu Leu Glu Arg Asp Ser Gly Leu Phe Pro Val Gly ** Ser Ser Asp Trp Ser Tyr Phe Ser Glu Ile Pro Gly Trp Ser His Tyr Glu Glu Val Ala Thr Gly Ala Thr Ser Ala Arg ** Arg GGG GGT CCT TAC CAT GAG GAG TTG ACG ACA GGG TCG ACA TCT TCG AGA GAT AGC 819 828 837 846 855 864 CCC CCA GGA ATG GTA CTC CTC AAC TGC TGT CCC AGC TGT AGA AGC TCT CTA TCG Pro Pro Gly Met Val Leu Leu Asn Cys Cys Pro Ser Cys Arg Ser Ser Leu Ser Pro Gln Glu Trp Tyr Ser Ser Thr Ala Val Pro Ala Val Glu Ala Leu Tyr Arg Pro Arg Asn Gly Thr Pro Gln Leu Leu Ser Gln Leu ** Lys Leu Ser Ile Gly

Ser *** Lys Ala Ile Lys Ser Ser Gln Gln Leu Val Ile Trp Pro Pro Val Pro Asn Ser Ser Gln Leu Lys Pro Leu Ser Ser Ser Phe Leu Gly Arg Leu Tyr Leu Ile Val Val Lys Cys Asn Gln Phe Val Ala Pro Ser Cys Asp Val Ser Thr CTC CTA ATG ATG AAA CGT TAA AAC CTT CTG ACG ACC TCT TGT TAG GTG CCT CCA 873 882 891 900 909 918 GAG GAT TAC TAC TTT GCA ATT TTG GAA GAC TGC TGG AGA ACA ATC CAC GGA GGT Glu Asp Tyr Tyr Phe Ala Ile Leu Glu Asp Cys Trp Arg Thr Ile His Gly Gly Arg Ile Thr Thr Leu Gln Phe Trp Lys Thr Ala Gly Glu Gln Ser Thr Glu Val Gly Leu Leu Leu Cys Asn Phe Gly Arg Leu Leu Glu Asn Asn Pro Arg Arg Tyr

Arg Leu Gly Ile Gln Leu Leu Pro Gly Val Arg His Gly Lys Gly Het Tyr Phe Gly Phe Ala Ser Lys Phe Cys His Val Trp Gly Thr Gly Lys Glu Trp Ile Phe Gly Ser Pro Arg Asn Ser Ala Thr Ser Gly Gly Gln Ala Arg Lys Gly Tyr Leu TGG GCT TCC GGC TAA ACT TCG TCA CCT GGG TGG GAC ACG GGA AAA GGG TAT ATT 927 936 945 954 963 972 ACC CGA AGG CCG ATT TGA AGC AGT GGA CCC ACC CTG TGC CCT TTT CCC ATA TAA Thr Arg Arg Pro Ile *** Ser Ser Gly Pro Thr Leu Cys Pro Phe Pro Ile *** Pro Glu Gly Arg Phe Glu Ala Val Asp Pro Pro Cys Ala Leu Phe Pro Tyr Lys Pro Lys Ala Asp Leu Lys Gln Trp/Thr His Pro Val Pro Phe Ser His Ile Lys

Leu Asn Ser Leu Arg Lys Gln ... Met Thr Ile Thr Lys Ile Lys Ile Tyr Ile Val Ser Asp Lys Lys Asn Asp Cys Arg Leu Pro Lys ... Lys ... Glu Ile Phe ... Gln Thr Lys Lys Thr Ile Val Asp Tyr His Asn Lys Asn Bal 990 1008 1017 1026

AAT AAA TTA CTG AGT CTT TTT TGT TAT CAC ATC GTA ATG GTT TTT ATT TTT ATT

Asn Lys Leu Leu Ser Leu Phe Cys Tyr His Ile Val Het Val Phe Ile Phe Ile Ile Asn Tyr ... Val Phe Phe Val Ile Thr Ser ... Trp Phe Leu Phe Leu Phe ... Ile Thr Glu Ser Phe Leu Leu Ser His Arg Asn Gly Phe Tyr Phe Tyr Ser

Lys Ser Pro Arg Glu Pro Tyr Ile Arg Gln Ile Thr (ys Leu Tyr Asp Val Lys Asn Leu Pro Asp Lys Leu Ile Phe Glu Arg Phe Gln Val Tyr Ile Thr Leu Arg Het Leu Thr Lys Ser Leu Asn Glu Ser Asn Tyr Het Phe Leu Gly GTA AAT CTC CCA GAA AGT CCT ATT TAA GAG ACT TAA CAT GTA TTT ATC AGT TGG 1035 1044 1053 1062 1071 1080 CAT TTA GAG GGT CTT TC: GGA TAA ATT CTC TGA ATT GTA CAT AAA TAG TCA ACC His Leu Glu Gly Leu Ser Gly *** Ile Leu *** Ile Val His Lys *** Ser Thr Ile *** Arg Val Phe Gln Asp Lys Phe Ser Glu Leu Tyr Ile Asn Ser Gln Pro Phe Arg Gly Ser Phe Arg Ile Asn Ser Leu Asn Cys Thr *** Ile Val Asn Leu

FIGURE 2 (continuation 3)

Gly Cys Leu Lys Pro Ser His Asn Cys Lys Pro Ala Cys Leu Gly Pro Arg His Val Val Tyr Asn Gln Ala Thr Thr Ala Asn Gln Leu Ald Tyr Gly Leu Gly Thr Trp Het Ile Lys Pro Gln Pro Gln Met Lys Ser Arg Het Ala Trp Ala Gln AAT GGT GTA TTA AAA CCC GAC ACC AAC GTA AAA CCT CGC GTA TCG GGT CCG GAC 1089 1098 1107 1116 1125 1134

TTA CCA CAT AAT TTT GGG CTG TGG TTG CAT TTT GGA GCG CAT AGC CCA GGC CTG Leu Pro His Asn Phe Gly Leu Trp Leu His Phe Gly Ala His Ser Pro Gly Leu Tyr His Ile Ile Leu Gly Cys Gly Cys Ile Leu Glu Arg Ile Ala Gln Ala Cys Thr Thr ** Phe Trp Ala Val Val Ala Phe Trp Ser Ala ** Pro Arg Pro Val

Ala Arg Cys Gln His Pro Tyr Lys Phe Pro Ala Val Ala Pro Lys Lys

His Glu Val Asn Thr His Thr Asn Leu His Leu Trp Leu Gln Asn Arg Lys Asn

Thr Ser Ser Met Pro Thr Pro Ile

ACA CGA GCT GTA ACC ACA CCC ATA AAT TTA CCT CGG TGT CGA CCA AAG AAA ATA

1143

1152

1161

1170

1179

1188

TGT GCT CGA CAT TGG TGT GGG TAT TTA AAT GGA GGC ACA GCT GGT TTC TTT TAT

Cys Ala Arg His Trp Cys Gly Tyr Leu Asn Gly Ala Thr Ala Gly Phe Phe Tyr

Val Leu Asp Ile Gly Val Gly Ile

Cys Ser Thr Leu Val Trp Val Phe Lys Trp Ser His Ser Trp Phe Leu Leu Leu

Lys Ala Pro Val Leu *** Asn Asn Pro Arg Ala Arg Thr Gln Pro His Leu Val Asn Pro Gln Phe Trp Asp Ile Thr Gln Asp Leu Glu Pro Lys Pro Thr Phe Tyr Ile Gln Ser Ser Gly Ile Leu Gln Lys Thr ** Ser Gln Asn Pro Pro Ser Thr ATA AAC CGA CCT TGG TTA GTT AAC AAA CCA GAT CGA GAC CAA ACC CCC ACT TCA 1197 1206 1215 1224 1233 1242
TAT TTG GCT GGA ACC AAT CAA TTG TTT GGT CTA GCT CTG GTT TGG GGG TGA AGT Tyr Leu Ala Gly Thr Asn Gln Leu Phe Gly Leu Ala Leu Val Trp Gly *** Ser Ile Trp Leu Glu Pro Ile Asn Cys Leu Val *** Leu Trp Phe Gly Gly Glu Val Phe Gly Trp Asn Gln Ser Ile Val Trp Ser Ser Gly Leu Gly Val Lys Tyr

Gln Leu Pro Leu Tyr Leu Ala Ala Lys His His Pro Pro Leu Leu Leu Tyr Arg Ser His Tyr Thr Phe Pro Gln Arg Ile Thr His Arg Ser Ser Tyr Asn Ile Gly Pro Thr Thr Pro Leu Pro Ser Gly Pro Thr Ala Pro Pro Thr Thr Leu TGG ACC TCA CCA TCC ATT TCC CGA CGG AAT ACC ACA CCG CCC TCC TCA TCA ATT 1251 1260 1269 1278 1287 1296 ACC TGG AGT GGT AGG TAA AGG GCT GCC TTA TGG TGT GGC GGG AGG AGT AGT TAA Thr Trp Ser Gly Arg Arg Ala Ala Leu Trp Cys Gly Gly Arg Ser Ser Pro Gly Val Val Gly Lys Gly Leu Pro Tyr Gly Val Ala Gly Gly Val Val Asn Leu Glu Trp Val Lys Gly/Cys Leu Het Val Trp Arg Glu Glu Leu Ile

Leu Pro *** Leu Gly Leu Gln His Leu Pro Asn Cys Leu Gln Cys Gly Leu Tyr Tyr Pro Asp Tyr Ala Leu Asn Thr Ser Pro Thr Val Phe Asn Ala Asp Leu Ile Ile Pro Thr Het Pro Trp Thy Pro Pro Pro Pro Pro Leu Thr Pro Het Trp Ser ATA TCC CCA GTA TCC GGT TCA ACC TCC CCC AAT GTT TCA ACC GTA GGT TCT 1305 1314 1323 1332 1341 1350 TAT AGG GGT CAT AGG CCA AGT TGG TGG AGG GGG TTA CAA AGT TGG CAT CCA AGA Tyr Arg Gly His Arg Pro Ser Trp Trp Arg Gly Leu Gln Ser Trp His Pro Arg Ile Gly Val Ile Gly Gln Val Gly Gly Gly Gly Tyr Lys Val Gly Ile Gln Asp Gly Ser ** Ala Lys Leu Val Glu Gly Val Thr Lys Leu Ala Ser Lys Ile

FIGURE 2 (continuation 4)

Cys Cys His Val Trp Cys Arg Lys Ser ** Leu His His Pro Arg Gln Pro Leu Val Val Thr Ser Gly Val Gly Arg Gln Asn Ser Thr Ile Pro Asp Arg Pro Tyr Leu Leu Leu Pro Gly Leu Val Glu Lys Ile Leu Pro Ser Pro Thr Glu Pro Thr ATT GTT GTC ACC TGG GTT GTG GAG AAA CTA ATC TCC ACT ACC CCA GAG ACC CCA 1359 1368 1377 1386 1395 1404
TAA CAA CAG TGG ACC CAA CAC CTC TTT GAT TAG AGG TGA TGG GGT CTC TGG GGT
*** Gln Gln Trp Thr Gln His Leu Phe Asp *** Arg *** Trp Gly Leu Trp Gly Asn Asn Ser Gly Pro Asn Thr Ser Leu Ile Arg Gly Asp Gly Val Ser Gly Val Thr Thr Val Asp Pro Thr Pro Leu *** Leu Glu Val Het Gly Ser Leu Gly ****

File *** Gly Lys *** Tyr Pro Leu Ile Pro Phe Thr Pro Thr Pro Pro Phe Glu Tyr Lys Ala Lys Arg Ile Arg Tyr Tyr Gln Phe Pro Leu Pro Leu Pro Phe Asn Het Asn Leu Arg Glu Leu Val Thr Thr Asn Ser Leu Tyr Pro Tyr Pro TTT TAA GTA TAA ATC GGA AAG ATT ATG CCA TCA TAA CCT TTC CAT CCC CAT CCC 1413 1422 1431 1440 1449 1458 AAA ATT CAT ATT TAG CCT TTC TAA TAC GGT AGT ATT GGA AAG GTA GGG GTA GGG Lys Ile His Ile *** Pro Phe *** Tyr Gly Ser Ile Gly Lys Val Gly Val Gly Lys Phe Ile Phe Ser Leu Ser Asn Thr Val Val Leu Glu Arg *** Gly *** Gly Asn Ser Tyr Leu Ala Phe Leu Ile Arg *** Tyr Trp Lys Gly Arg Gly

Gln His Arg Arg Leu Pro Pro Pro Val Pro Arg His Gln Ile Glu Ala Arg Asn Thr Gly Gly Ser Pro Pro Leu Phe Gln Gly Ile Asn Phe Arg Leu Glu Asn Thr Pro Ala Ala Gln Pro Pro Ser Ser Ser Ala Ser Thr Ser Asp Ser Thr CCA ACC ACG GCG GAC TCC CCC CCT CCT TGA CCG GCT ACA ACT TAG AGT CGA GCA 1467 1476 1485 1494 1503 1512 GGT TGG TGC CGC CTG AGG GGG GGA GGA ACT GGC CGA TGT TGA ATC TCA GCT CGT Gly Trp Cys Arg Leu Arg Gly Gly Gly Thr Gly Arg Cys Ile Ser Ala Arg Val Gly Ala Ala Gly Gly Gly Glu Glu Leu Ala Asp Val Glu Ser Gln Leu Val Leu Val Pro Pro Glu Gly Gly Arg Asn Trp Pro Het Leu Asn Leu Ser Ser Leu

Cys Glu Leu Ile Ala Ala Leu Thr Arg Arg Lys His His Thr Cys Ile Arg val Asn Trp Ser Pro Gln Ser His Gly Gly Arg Ile Thr Leu Val Phe Glu Arg Leu Met Gly Leu His Ser Arg Thr Asp Glu Glu Pro Ser Tyr Leu Asn Glu ATT GTA AGG TTC TAC CGA CGC TCA CAG GAG GAG AAT ACC ACT CAT GTT TAA GAG 1521 1530 1539 1548 1557 1566

TAA CAT TCC AAG ATG GCT GCG AGT GTC CTC CTC TTA TGG TGA GTA CAA ATT CTC His Ser Lys Met Ala Ala Ser Val Leu Leu Trp Vol Gln Ile Leu Asn Ile Pro Arg Trp Leu Arg Val Ser Ser Ser Tyr Gly Glu Tyr Lys Phe Ser Thr Phe Gln Asp Gly Cys/Glu Cys Pro Pro Leu Met Val Ser Thr Asn Ser Leu

Phe Pro Pro Phe Gln Leu Tyr Gly Asp Lys Pro Ala Met Gln Leu Pro Lys Gln Ser Leu Arg Ser Asn Phe Ile Gly Thr Lys Arg Arg Trp Arg Tyr Arg Asn Arg Leu Phe Ala Pro Ile Ser Ser Val Arg Arg Glu Ala Gly Asp Thr Val Thr Glu ATC TTT CCG CCC TTA ACT TCT ATG GGC AGA AAG CCG CGG TAG ACA TTG CCA AAG 1575 1584 1593 1602 1611 1620 TAG AAA GGC GGG AAT TGA AGA TAC CCG TCT TTC GGC GCC ATC TGT AAC GGT TTC Lys Gly Gly Asn *** Arg Tyr Pro Ser Phe Gly Ala Ile Cys Asn Gly Phe Arg Lys Ala Gly Ile Glu Asp Thr Arg Leu Ser Ala Pro Ser Val Thr Val Ser Glu Arg Arg Glu Leu Lys Ile Pro Val Phe Arg Arg His Leu *** Arg Phe Leu

FIGURE 2 (continuation 5)

Leu Arg Pro Thr Gly Phe Ile Thr Lys Glu Pro Pro His Lys Trp Ser Pro Gln Phe Ala Pro His Val Leu Tyr Pro Arg Arg Arg Leu Ile Asn Gly Leu His Ser Ser Pro Pro Thr Tyr Trp Ile His Asp Glu Gly Ser Ser Thr Glu Leu Ile Ala ACT TCC GCC CCA CAT GGT TTA TAC CAG AAG AGG CCT CCT ACA AAG GTT CTA CCG 1629 1638 1647 1656 1665 1674
TGA AGG CGG GGT GTA CCA AAT ATG GTC TTC TCC GGA GGA TGT TTC CAA GAT GGC Arg Arg Gly Val Pro Asn Met Val Phe Ser Gly Gly Cys Phe Gln Asp Gly Glu Gly Gly Val Tyr Gln Ile Trp Ser Ser Pro Glu Asp Val Ser Lys Met Ala Lys Ala Gly Cys Thr Lys Tyr Gly Leu Leu Arg Arg Het Phe Pro Arg Trp Leu

Pro Pro Pro Asp Thr Lys Gln Pro Leu Ala Glu Lys Ala Val Asp Asp ... Leu Arg Pro Arg Thr Arg Arg Arg Arg Arg Arg Pro Trp Thr Het Arg Tyr Ala Pro Ala Pro Gly Asp Glu Ala Thr Val Gly Gly Gln Gly Arg ... Gly Ile ACG CCC CCG CCC AGG CAG AAG ACG CCA TTG CGG AGG AAC CGG TGC AGT AGG ATA 1683 1692 1701 1710 1719 1728 TGC GGG GGC GGG TCC GTC TTC TGC GGT AAC GGC TCC TTG GCC ACG TCA TCC TAT Cys Gly Gly Gly Ser Val Phe Cys Gly Asn Ala Ser Leu Ala Thr Ser Ser Tyr Ala Gly Ala Gly Pro Ser Ser Ala Val Thr Pro Pro Trp Pro Arg His Pro Ile Arg Gly Arg Val Arg Leu Leu Arg ... Arg Leu Leu Gly His Val Ile Leu ...

Leu Ser Leu Leu Ala Ser Ser Tyr Tyr
Phe His Phe Phe His Ala Ala Thr Thr Asn
Phe Thr Phe Ser Thr Arg Gln Gln Leu I'le

TTT TCA CTT TCT TCA CGC GAC GAC ATC ATA A 5.

1737 1746 1755

AAA AGT GAA AGA AGT GCG CTG CTG TAG TAT T 3.

Lys Ser Glu Arg Ser Ala Leu Leu . Tyr
Lys Val Lys Glu Val Arg Cys Cys Ser Ile
Lys . Lys Lys Cys Ala Ala Val Val

FIGURE 2 (continuation 6).

3	_/	2		
7	_	4	0	

	10 70	
circopormank circopormeeh	1 ACCAGGGCAC PICGGCAGCG GCAGCACGTC GGCAGCGTCA GTGAAAATGC	SO
circopordfp	1 SECAGEGEAE FIEGGEAGEG SCAGEACETE GGEAGEGTEA GTGAAAATGE	50 50
Circopormank	51 AAGGAAGAA 3AGGGGGGGG AAGGGGGGGGGGGGGGG	
Circopormech		100
circopordfp	SI CAAGCAAGAA CAGCGGCCCG CAACCCCATA GGAGGTGGGT STTCACCCTT	100 100
	110 120 130 140 150	
circopormank Circopormeeh	TOT PATRATECTT COAGGAGGA GAAAAACAAA ATACGGGAGC TTCCAATCTC	150
circopordfp	101 SATAATCCTT CCGAGGAGGA GAAAAACAAA STACGGGAGC FTCCAATCTC	150 150
	160 170 / 180 190 200	
Circopormank	151 CHITTIGAT PATTITIGITI GEGGAGAGGA AGGTTTGGAA GAGGGTAGAA	200
circopormeeh	151 CHERTIGAT PATETTIGETT GEGGAGAGGA AGGTTTGGAA GAGGGTAGAA	200
circopordfp	151 STREETGAT PATRITUGETT STEEGE SAGGA SGGTAGAA	200
•	210 220 / 230 240 250	
circopormank	201 STEECACCT CCAGGGGTTT GCTAATTTTG CTAAGAAGCA GACTTTTAAC	250
circopormeeh circopordfp	201 SICCICACCI SCAGGGGTTT GCGAATTTTG STAAGAAGCA GACTTTTAAC	250
cti copoi ai p	201 STECTCACCT SCAGGGGTTT GCGAATTTTG STAAGAAGCA GACTTTTAAC	250
	260 270/ 280 290 300	
circopormank	251 AAGGIGAAGT AGTATITITGG FGCCCGCTGC CACATCGAGA AAGCGAAAGG	300
circopormeeh circopordfp	251 SAGGTGAAGT AGTATITITGG FGCCCGCTGC CACATCGAGA SAGCGAAAGG 251 SAGGTGAAGT AGTATITITGG FGCCCGCTGC CACATCGAGA SAGCGAAAGG	300
	251 LAGGIGAAGT SGTATTITGG TGCCCGCTGC CACATCGAGA SAGCGAAAGG	300
	310 /320 330 340 350	
circopormank circopormeeh	301 JACCGACCAG CAGAATAAAG JATACTGCAG FAAAGAAGGC CACATACTTA	350
circopordfp	301 MACCGACCAG CAGAATAAAG MATACTGCAG TAAAGAAGGC CACATACTTA 301 MACCGACCAG CAGAATAAAG MATACTGCAG TAAAGAAGGC CACATACTTA	350
		350
circopormank	360 / 370 380 390 400	
Circopormeeh	351 ICGAGTGTGG AGCTCCGCGG AACCAGGGGA AGCGCAGCGA CCTGTCTACT	400
circopordfp	351 ICGAGTGTGG AGCTCCGCGG AACCAGGGGA AGCGCAGCGA CCTGTCTACT 351 ICGAGTGTGG AGCTCCGCGG AACCAGGGGA AGCGCAGCGA CCTGTCTACT	400
		400
Circopormank	410/ 420 430 440 450 401 विकास स्वित्ता स्वापन	
circopormeeh	ACT TO COLOR OF THE PROPERTY O	450
circopordfp	401 GETGTGAGTA SCETTTTGGA GAEGGGGTET TTGGTGAETG FAGEEGAGEA	450 450
circopormank	/460 470 480 490 500	
circopormeeh	451 STICCETGTA ACGTATGTGA GAAATTTCCG EGGGCTGGCT GAACTTTTGA 451 STICCETGTA ACGTATGTGA GAAATTTCCG EGGGCTGGCT GAACTTTTGA	500
circopordfp	451 STUSCHGIA ACGIATGIGA SAAATTICCG CGGGGGGGGGT GAACTTITGA 451 STUSCHGTA ACGIATGIGA SAAATTICCG CGGGCTGGCT GAACTTITGA	500 500
		200
circopormank	510 520 530 540 550 501 अनुस्तित्त्ति त्रश्तिकात्त्रत् स्वत्त्तात्राम् स्वयंत्रत्यस्य स्वरंभवत्त्रत्	.
circopormeeh	CO. MICHELL STREET	\$50 \$50
circopordfp	501 SAGTGAGEGG SAAGATGEAG SAGEGTGATT SGAAGAEAGE FGTAEAEGTE	550 550
		٠ .
circopormank	560 570 580 590 600 551 अर्थानन्त्व त्ववववन्तान त्ववक्षत्त्व क्षत्रत्वत्वव त्रक्षक्ष्यक्त्व	
circopormeeh	551 TAGTGGGCC FGCCCGGTTG TGGGAAGAGC FAGTGGGCCC GTAATTTTGC	600 600
circopordfp	\$51 TAGTGGGCC EGCCCGGTTG TGGGAAGAGC CAGTGGGCCC STAATTTTGC	- 600
		-

circopormank circopormech circopordfp	601 TGAGCCTAGG GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG 601 TGAGCCTAGG GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG 601 TGAGCCTAGG GACACCTACT GGAAGCCTAG TAGAAATAAG TGGTGGGATG	650 650 650
circopormank circopormech circopordfp	651 GATATCATGG AGAAGAAGTI STIGTTITGG ATGATTTITA IGGGTGGTTA 651 GATATCATGG AGAAGAAGTI STIGTTITGG ATGATTTITA IGGGTGGTTA 651 GATATCATGG AGAAGAAGTI STIGTTITGG ATGATTTITA IGGGTGGTTA	700 700 700
circopormank circopormeeh circopordfp	701 SCHIGGGATG ATCTACTGAG ACTGTGTGAC SCGTATCCAT TGACTGTAGA ACTGTGTGAC SCGTATCCAT TGACTGTAGA ACTGTGTGAC SCGTATCCAT TGACTGTAGA ACTGTGTGAC SCGTATCCAT TGACTGTAGA	750 750 750
circopormank circopormeeh circopordfp	751 JACTANAGGE AGTACTOTIC STITUTIOGE TEGENGRATI FIGATIACCA 751 JACTANAGGE AGTACTOTIC STITUTIOGE ECGENGRATI FIGATIACCA 751 JACTANAGGE AGTACTOTIC STITUTIOGE ECGENGRATI FIGATIACCA 751 JACTANAGGE AGTACTOTIC STITUTIOGE ECGENGRATI FIGATIACCA	800 800 800
circopormank circopormeeh circopordfp	801 GCAATCAGGC GCCCCAGGAA GGTACTCCT CAACTGCTGT GCCAGCTGTA 801 GCAATCAGGC GCCCCAGGAA GGTACTCCT CAACTGCTGT GCCAGCTGTA 801 GCAATCAGGC GCCCCAGGAA GGTACTCCT CAACTGCTGT GCCAGCTGTA 601 GCAATCAGGC GCCCCAGGAA GGTACTCCT CAACTGCTGT GCCAGCTGTA	850 850 850
circopormank circopormeeh circopordfp	851 GAAGGIGIGI ATCGGAGGAT FACTACTITIC CAATTITIGGA AGACTGCTGG 851 GAAGGIGIGI ATCGGAGGAT FACTACTITIC CAATTITIGGA AGACTGCTGG 851 GAAGGIGIGI ATCGGAGGAT FACTACTITIC CAATTITIGGA AGACTGCTGG	900 900 900
circopormank circopormeeh circopordfp	901 AGAACAATICA ACGGAGGTAC CGAAGGCCG ATTTGAAGCA ATGGACCCAC STOTTGAAGCA ATGGACCCAC SCGAAGGCCG ATTTGAAGCA ATGGACCCAC SCGAAGGCCG ATTTGAAGCA ATGGACCCAC SCGAAGGCCG ATTTGAAGCA ATGGACCCAC STOTTGAAGCA ATGGACCCAC	950 950 950
circopormank circopormeeh circopordfp	951 STUTIGEST TECCCATAT SAAATAAATT SCIGAGIGIT TITIGITATC	1000 1000 1000
circopormank circopormeeh circopordfp	1001 SCATEGIAAT SCHEFFATT FITATTIATT FAGAGGGTCT TTTAGGATAA	1050 1050 1050
circopormank circopormeeh circopordfp	1051 STICTICIONE PIGNACATAN STAGTCANEC FTACCACATA STITTGGGGT	1100 1100 1100
circopormank circopormeeh circopordfp	1101 Inder Made Andre Stage Stage Setting Concentration	1150 1150 1150
circopormank circopormeeh circopordfp	1151 GIGGIATITI SAATGGAGCC SCAGGIGGTI ICTITATIA HIGGGIGGA 1151 GIGGIATITI SAATGGAGCC SCAGGIGGTI ICTITATIA HIGGGIGGA 1151 GIGGIATITI SAATGGAGCC SCAGGIGGTI ICTITATIA HIGGGIGGA	1200 1200 1200

FIGURE 3 (continuation 1)

		11/26	/ .	
circopormank circopormeeh circopordfp	1301 ACCAATCAAT (GT	1220 1230 HIGGICE AGGICAGGIT HIGGICE AGGICAGGIT HIGGICE AGGICAGGIT	FGGGGGTGAA STAC	1250 CTGGAG 1250 CTGGAG 1250 CTGGAG 1250
Circopormank Circopormeeh circopordfp	1251 FGGTAGGTAA AGG 1251 FGGTAGGTAA AGG	1270 1280 GETGCET FATGGTGTGG GETGCET FATGGTGTGG	GGGAGGAGT AGTT	1300 AATATA 1300 AATATA 1300 AATATA 1300
circopormank circopormeeh circopordfp	1301 GGGGTCATAG GCC	1320 1330 AAGTIGG FGGAGGGGGT AAGTIGG FGGAGGGGGT AAGTIGG FGGAGGGGGT	FACAAAGTTG GCAT	1350 CCAAGA 1350 CCAAGA 1350 CCAAGA 1350
circopormank circopormeeh circopordfp	1351 FAACAACAGT GGAG 1351 FAACAACAGT GGAG 1351 FAACAACAGT GGAG	CCAACA CICITICAT CCAACA CICITICAT CCAACA CICITICAT	FAGAGGTGAT GGGG	1400 IGIGIG 1400 IGIGIG 1400 IGIGIG 1400
circopormank circopormeeh circopordfp	1401 GGGTAAAATT CATA	ATTAGE STRETATA	EGGTAGTATT GGAA	1450 3GGIAG 1450 3GGIAG 1450 3GGIAG 1450
circopormank circopormeeh circopordfp	1451 GGGTAGGGGG FTGG		GAGGAACTGG CCGAT	1500 GHGA 1500 GHGA 1500 GHGA 1500
circopormank circopormeeh circopordfp	1501 STETGAGGT STEA	ACATIC CAAGATGGCT	GCGAGTATCC CCCT	1550 11A1G 1550 11A1G 1550 11A1G 1550
circopormank circopormeeh circopordfp	1551 STGATTACAA STTC	TUTAGA AAGGCGGGAA	FTGAAGATAC CCGTC	1600 1600 1600 1650
circopormank circopormeeh circopordfp	1601 SCGCCATCTG FAAC	GGTTTC FGAAGGCGGG GGTTTC FGAAGGCGGG	GTGTGCCAAA FATG	1650
circopormank circopormeeh circopordfp	1651 CCGGAGGAT/SITE	CCAAGA FGGCTGCGGG CCAAGA FGGCTGCGGG CCAAGA FGGCTGCGGG	GGCGGGTCCT FCTT	
circopormank circopormeeh circopordfp	1701 PAREGETTE FIGO 1701 PAREGETTE FIGO	ATCCTATAA ATCCTATAA ATCCTATAA 1770 1780	AAGTGAAAGA AGTG	1750 1750 1750 1750
circopormank circopormeeh circopordfp		• • • • • • • • • • • • • • • • • • • •		1800

FIGURE 3 (continuation 2)

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- •		10	20	30	40	50	
circopormank	7	MPSKKSGPQP	HKRWVFTLNN	PSEEEKNKIR	ELPISLEOYE	VCGEEGLEEG	50
circopormeeh	1	MPSKKSGPQP	HKRWVFTLNN	PSEEEKNKIR	LPISLFOYF	VCGEEGLEEG	50
circopordfp[+	MPSKKSGPQP	HKRWVFTLNN	PSEEEKNKIR	ELPISLEDYE	VCGEEGLEEG	50
		60	70	80	90	100	
circopormank	51	REAHLOGEAN	FAKKOTENKY	KWYFGARCHI	EKAKGTOOON	KEYGSKEGHT	100
circopormeeh	51	RTPHLOGFAN	FAKKOTFNKV	KWYFGARCHI	EKAKGTOOON	KEYCSKEGHT	100
circopordfp[51	RTPHLOGFAN	FAKKOTFNKV	KWYFGARCHI	EKAKGTOOON	KEYCSKEGHT	100
				7			
		110	120		140	150	
circopormank	101	LIECGAPRNQ	GKRSDLSTAV	STLLETGSLV	TVAEQFPVTY	VRNFRGLAEL	150
circopormeeh	101	LIECGAPRNQ	GKRSDLSTAV	STLLETGSLV	TVAEQFPVTY	VRNFRGLAEL	150
circopordfp[101	LIECGAPRNO	GKRSDLSTAV	STLLETGSLV	TVAEOFPVTY	VRNFRGLAEL	150
				/ ·			
•		160	170	/ 180	190	Z00	
circopormank	151	LKYSGKMQQR	DIAKLWAHATA	GPPGCGKSQTI	ARNFAEPEDT	YVKPSRNKVITV	200
circopormeeh	151	LKVSGKMQQR	DWKTAVHVIV	GPPGCGKSQN	ARNFAEPROT	TIKPSRNKTT!	200
circopordfp[151	LKVSGKNOOR	DUKTAVHVIV	GPPGCGKSON	ARNFAEPROT	YVKPSRNKYTY	200
•							
		210	220	/ 230	240	250	
circopormank	201	DGYHGEEVVV	LODFYGWLPW		PLTVETKGGT	VPFLARSILI	250
circopormeeh	201	DGYHGEEVVV	LODFYGWLPW	DOLLRLCORY	PLTVETKGGT	VPFLARSILI	250
circopordfp[201	DGYHGEEVVV	DDFYGWLPW	JOLEKI CORY	जनग्र सम्	VEEL AREA	250
			/				
		260	2/70	280	290	300	200
circopormank	251		STAVPAVEA	YRRITTLOF	YKTAGEOSTE	VPEGREEAVD	300
circopormeeh		TSNQAPQEWY			YKTAGEQSTE	VPEGRFEAVO	300
circopordfp[251	TSNOAPOEWY	STAVPAVEA	LYRRITTLOF	YKTAGEOSTE	UPEGRFEAVD	300
		310	320	220	340	350	
<i>a:</i>	201			330	- ·		350
circopormank	301	PPCALFPYKI			• • • • • • • •		350
circopormeeh	301	PPCALFPYKI			• • • • • • • • •		350
circopordfp[301		/	• • • • • • • •	• • • • • • • • •	• • • • • • • •	2.20

13	/	2	6
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		10	20	30	1 40	50	
circopormank	1	MTYPRRRYRR	RRTRPRSHLG		MPAFRNRYRW	RRKTGIFNER	50
circopormeeh circopordfp[1	ATWPRRRYRR ATWPRRRYRR	RRTRPRSHLG RRTRPRSHLG		PAFRNRYRW	RRKTGIFNSR	50
ctreoporatpt		and the state of t	HATRIPR SHE U	MILRRRPYLV	HPAFRNRYRW	RRKTGIENSR	50
	·	60	70	80	90	100	
circopormank	51	SK SEVICER	3GYSOPSW IV	N RENTGOE	90 IPPSGGTNPI	PLPFOYYRIR	100
circopormeeh	51	STEVETK	GYSOPSWNV	TOP IN SAME	PPSGGTNPL	PLPFOYYRIR	100
circopordfp[51	SEFVETIR	GGISOPSWNV	NE RENIGOE	PPSGGTNPL	PLPFOYYRIR	100
				7			
•		110	120		140	150	
circopormank	101	KAKYEFYPRO	TISNERGVG	STVVILDANF	VTPSTNLAYD	PYINYSSRHT	150
circopormeeh	101	KAKYEFYPRD	ITSNORGVG	STVVILDANF	VTPSTNLAYD	YINYSSRHT	150
circopordfp[101	KAKYEFYPRO	PITSNORGVG	STATEDANE	VTPSTNLAYD	PYTAYSSRIFT	150
		160	170	/ 180	100	300	
circopormank	151	TROPFTYHER	VETEKPELOO	TE YEHENNK	RNOTHER TANK	200	700
circopormeeh	151	TROPFTYHSR	YFTPKPELDO	TIOWFHPNNK	RNOLWLHLNT	TINVENTALA	200 200
circopordfp[151	TROPFTYHSR	YFTPKPELDO	FIDWEDPNNK	RNOLWLHLNT	HTNVEHTGLG	200
			•				200
		210	220	/ 230	240	250	
circopormank	201.	PALQNAATAQ	IYVVRLTIYV	QFREFILKOP	ENK-		250
circopormeeh	201	YAL QNAATAQ		DEREFILKOP		• • • • • • • • •	250
circopordfp[201	YALONATIAO	HYURLHYU	DEREFILKOP	I IE•	• • • • • • • •	250
					•		
			/	_			•
•			FIGURI	E 5	•		•
•		·	FIGURI	E 5	•		•
			FIGURI	E 5	•		•
•		10	FIGURI 20	E 5	40	50	
circopormank	1	10 MISIPPLIST		30	40 FGRAHYDVYS	SO SLPETLINE	S 0
circopormeeh	1	MISIPPLIST MISIPPLIST	RLPVGVARLS RLPVGVPRLS	30 KITGPLALPT KITGPLALPT	TGRAHYDVYS TGRAHYDVYS	CLPITLLHLP	S 0 S 0
_	1 1 1	MISIPPLIST	RLPVGVARLS	30 KITGPLALPT KITGPLALPT	TGRAHYDVYS	CLPITLLHLP	
circopormeeh	1 1 1	WISTPPLIST WISTPPLIST WISTPPLIST	RLPVGVARLS RLPVGVPRLS RLPVGVPRLS	30 KITGPLALPT KITGPLALPT KITGPLALPT	TGRAHYDVYS TGRAHYDVYS TGRAHYDVYS	CLPITLLHLP	50
circopormeeh circopordfp[1 1 1	MISTPPLIST WISTPPLIST WISTPPLIST	RLPYGVARLS RLPYGVPRLS RLPYGVPRLS	30 KITGPLALPT KITGPLALPT KITGPLALPT	TGRAHYDVYS TGRAHYDVYS TGRAHYDVYS	IPITLUHUP IPITLUHUP IPITLUHUP	50 50
circopormeeh circopordfp[circopormank	1 1 1 51	MISTPPLIST WISTPPLIST WISTPPLIST 60 AHFQKFSQPA	20 RLPYGVARLS RLPYGVPRLS RLPYGVPRLS 70 FISHIRYREL	30 KITGPLALPT KITGPLALPT KITGPLALPT 80 LGYSHQRPRL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV	SLPITLLHLP SLPITLLHLP 100 SALPLVPRSS	50 50 100
circopormeeh circopordfp[circopormank circopormeeh	51	MISTPPLIST MISTPPLIST MISTPPLIST MISTPPLIST 60 AHFQKFSQPA AHFQKFSQPA	20 RLPYGVARLS RLPYGVPRLS RLPYGVPRLS 70 FISHIRYREL FISHIRYREL	30 KITGPLALPT KITGPLALPT KITGPLALPT ETGPLALPT 80 LGYSHQRPRL LGYSHQRPRL	TGRAHYDVYS TGRAHYDVYS TGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV	100 SALPLVPRSS SALPLVPRSS	50 50 100 100
circopormeeh circopordfp[circopormank		MISTPPLIST WISTPPLIST WISTPPLIST 60 AHFQKFSQPA	20 RLPYGVARLS RLPYGVPRLS RLPYGVPRLS 70 FISHIRYREL	30 KITGPLALPT KITGPLALPT KITGPLALPT 80 LGYSHQRPRL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV	SLPITLLHLP SLPITLLHLP 100 SALPLVPRSS	50 50 100
circopormeeh circopordfp[circopormank circopormeeh	51	MISTPPLIST MISTPPLIST MISTPPLIST MISTPPLIST 60 AHFQKFSQPA AHFQKFSQPA	20 RLPYGVARLS RLPYGVPRLS RLPYGVPRLS 70 FISHIRYREL FISHIRYREL	30 KITGPLALPT KITGPLALPT KITGPLALPT ETGPLALPT 80 LGYSHQRPRL LGYSHQRPRL	TGRAHYDVYS TGRAHYDVYS TGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV	100 SALPLVPRSS SALPLVPRSS	50 50 100 100
circopormeeh circopordfp[circopormank circopormeeh	51	MISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST AHFOKESOPA AHFOKESOPA AHFOKESOPA AHFOKESOPA	RLPVGVARLS RLPVGVPRLS RLPVGVPRLS RLPVGVPRLS 70 RSHIRVREL RSHIRVREL RSHIRVREL RSHIRVREL	30 KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHORPRL GYSHORPRL GYSHORPRL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 OKGTHSSROV OKGTHSSROV OKGTHSSROV	100 SALPLVPRSS AALPLVPRSS AALPLVPRSS	50 50 100 100
circopormeeh circopormank circopormeeh circopordfp[circopormank circopormank	51 51	WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST 60 AHFOKESOPA AHFOKESOPA AHFOKESOPA 110 FLDKYVAFFT	RLPVGVARLS RLPVGVPRLS RLPVGVPRLS RLPVGVPRLS 70 FISHIRYREL FISHIRYREL FISHIRYREL FISHIRYREL FISHIRYREL FISHIRYREL	30 KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 OKGTHSSROV OKGTHSSROV OKGTHSSROV 140 KIPLHLVKSL	100 SALPLVPRSS SALPLVPRSS SALPLVPRSS	50 50 100 100 150 150
circopormeeh circopormank circopormeeh circopordfp[51 51 101 101	WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST 60 AHFOKFSOPA AHFOKFSOPA AHFOKFSOPA 110 FLDKYVAFFT FLDKYVAFFT	RLPVGVARLS RLPVGVPRLS RLPVGVPRLS RLPVGVPRLS RLPVGVPRLS RLPVGVPRLS RVFEL RVREL	STGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL TRFLOVAAGT	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV	100 SALPLVPRSS SALPLVPRSS ALPLVPRSS LSKISKPLE LSKIRKPLE	50 50 100 100 150
circopormeeh circopormank circopormeeh circopordfp[circopormank circopormank	51 51 101 101	MISTPPLIST	RLPYGVARLS RLPYGVPRLS	30 KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL TRFLOVAAGT FRFLOVAAGT FRFLOVAAGT	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV	LPITLLHLP LPITLLHLP LPITLLHLP 100 SALPLVPRSS ALPLVPRSS ALPLVPRSS LSKISKPLE LSKIRKPLE LSKIRKPLE LSKIRKPLE	50 50 100 100 150 150
circopormeeh circopormank circopormeeh circopordfp[circopormank circopormeeh circopormeeh circopordfp[51 51 101 101 101	WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST WISTPPLIST 60 WHEOKESOPA WHEOKESOPA 110 FLDKYVAFFT FLDKYVAFFT FLDKYVAFFT FLDKYVAFFT FLDKYVAFFT	RIPVGVARLS RIPVGVPRLS	SO KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 OKGTHSSROV OKGTHSSROV OKGTHSSROV 140 KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL	100 SALPLVPRSS SALPLVPRSS SALPLVPRSS LSKIKKPLE LSKIRKPLE LSKIRKPLE	50 50 100 100 150 150
circopormeeh circopormank circopormeeh circopordfp[circopormank circopormeeh circopormeeh circopordfp[51 51 101 101 101	WISTPPLIST WISTPPLIST WISTPPLIST 60 WHEOKESOPA WHEOKESOPA WHEOKESOPA 110 FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT FLOKYVAFFT	RLPVGVARLS RLPVGVPRLS RVFFILLVGS	STEPLALPT KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL 130 FRFLDVAAGT FRFLDVAAGT FRFLDVAAGT	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV 140 KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL 190 LGRIJKGEH	100 SALPLVPRSS SALPLVPRSS SALPLVPRSS SALPLVPRSS LSKISKPLE LSKIRKPLE LSKIRKPLE LSKIRKPLE LSKIRKPLE LSKIRKPLE	50 50 100 100 150 150 150
circopormeeh circopormank circopormeeh circopormank circopormeeh circopormeeh circopormeeh circopormeeh	51 51 101 101 101 151	VISTPPLIST VISTPPLIST VISTPPLIST VISTPPLIST 60 AHFOKESOPA AHFOKESOPA AHFOKESOPA TLOKYVAFFT VLOKYVAFFT VLOKYVAFFT VRSSTLFOTF VRSSTLFOTF	RIPYGYARLS RIPYGYPRUS RIPYGYPRUS RIPYGYPRUS RIPYGYPRUS RIPYREL RISHIRYREL RIS	30 KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT JWKLPYFVFL DWKLPYFVFL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV	100 SALPLVPRSS SALPLVP	50 50 100 100 150 150 150 200 200
circopormeeh circopormank circopormeeh circopordfp[circopormank circopormeeh circopormeeh circopordfp[51 51 101 101 101 151	VISTPPLIST VISTPPLIST VISTPPLIST VISTPPLIST 60 AHFOKESOPA AHFOKESOPA AHFOKESOPA TLOKYVAFFT VLOKYVAFFT VLOKYVAFFT VRSSTLFOTF VRSSTLFOTF	RIPYGYARLS RIPYGYPRUS RIPYGYPRUS RIPYGYPRUS RIPYGYPRUS RIPYREL RISHIRYREL RIS	30 KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT JWKLPYFVFL DWKLPYFVFL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV 140 KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL 190 LGRIJKGEH	100 SALPLVPRSS SALPLVP	50 50 100 100 150 150 150
circopormeeh circopormank circopormeeh circopormank circopormeeh circopormeeh circopormeeh circopormeeh	51 51 101 101 101 151	VISTPPLIST VISTPPLIST VISTPPLIST VISTPPLIST 60 AHFOKESOPA AHFOKESOPA AHFOKESOPA TLOKYVAFFT VLOKYVAFFT VLOKYVAFFT VRSSTLFOTF VRSSTLFOTF	ILPYGYARLS REPYGYPRUS REPYGYPRUS REPYGYPRUS REPYGYPRUS REPYREL	ATTGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL TRFLOVAAGT FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT JWKLPYFVFL DWKLPYFVFL DWKLPYFVFL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV 140 LIPLHLVKSL KIPLHLVKSL KIPLHLVKSL LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH	100 GALPLVPRSS GALPLE GALPLVPRSS	50 50 100 100 150 150 150 200 200
circopormeeh circopormank circopormank circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopordfp[51 51 101 101 101 151 151	VISTPPLIST VISTPPLIST VISTPPLIST 60 WHEOKESOPA WHEOKESOPA WHEOKESOPA 110 VIDKYVAFFT VILDKYVAFFT	RIPYGVARIS RIPYGVPRIS RIPYGVPRIS RIPYGVPRIS RIPYREL RISHIRYREL RISHIRYREL RISHIRYREL RISHIRYRE R	30 KITGPLALPT KITGPLALPT KITGPLALPT KITGPLALPT 80 GYSHQRPRL GYSHQRPRL GYSHQRPRL GYSHQRPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT PRFLOVAAGT 180 DWKLPYFVFL DWKLPYFVFL DWKLPYFVFL	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 IKGTHSSROV IKGTHSSROV IKGTHSSROV ILGRIHVKSL KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH	IPITLLHIP IPITLLHIP IPITLLHIP IPITLLHIP IOO SALPLVPRSS	50 50 100 100 150 150 150 200 200 200
circopormeeh circopormank circopormank circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh	51 51 101 101 101 151 151 201 201	VISTPPLIST VISTPPLIST ANTERNATION ANTERNATION ANTERNATION LDKYVAFFT LDKYVAFFT LDKYVAFFT VRSSTLFOTF VRSSTLFOTF VRSSTLFOTF VRSSTLFOTF ANTERNATION ANTER	RIPYGVARLS RIPYGVPRLS	SO KITGPLALPT KITGPLALPT KITGPLALPT STGPLALPT STGPLALPT SO GYSHORPRL GYSHORPRL GYSHORPRL GYSHORPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT 180 DWKLPYFVFL DWKLPYFVFL DWKLPYFVFL 236	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV QKGTHSSRQV 140 LGRILKGEH LGRILKGEH LGRILKGEH 240	IPITLUILP IPITLUILP IPITLUILP IPITLUILP IOO SALPLVPRSS	50 50 100 100 150 150 200 200 200 250 250
circopormeeh circopormank circopormank circopormeeh circopormeeh circopormeeh circopormeeh circopormeeh circopordfp[51 51 101 101 101 151 151 201 201	VISTPPLIST VISTPPLIST ANTERNATION ANTERNATION ANTERNATION LDKYVAFFT LDKYVAFFT LDKYVAFFT VRSSTLFOTF VRSSTLFOTF VRSSTLFOTF VRSSTLFOTF ANTERNATION ANTER	RIPYGVARLS RIPYGVPRLS	SO KITGPLALPT KITGPLALPT KITGPLALPT STGPLALPT STGPLALPT SO GYSHORPRL GYSHORPRL GYSHORPRL GYSHORPRL 130 FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT FRFLOVAAGT 180 DWKLPYFVFL DWKLPYFVFL DWKLPYFVFL 236	IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS IGRAHYDVYS 90 IKGTHSSROV IKGTHSSROV IKGTHSSROV ILGRIHVKSL KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL KIPLHLVKSL LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH LIGRIIKGEH	IPITLUILP IPITLUILP IPITLUILP IPITLUILP IOO SALPLVPRSS	50 50 100 100 150 150 150 200 200 200

FIGURE 6

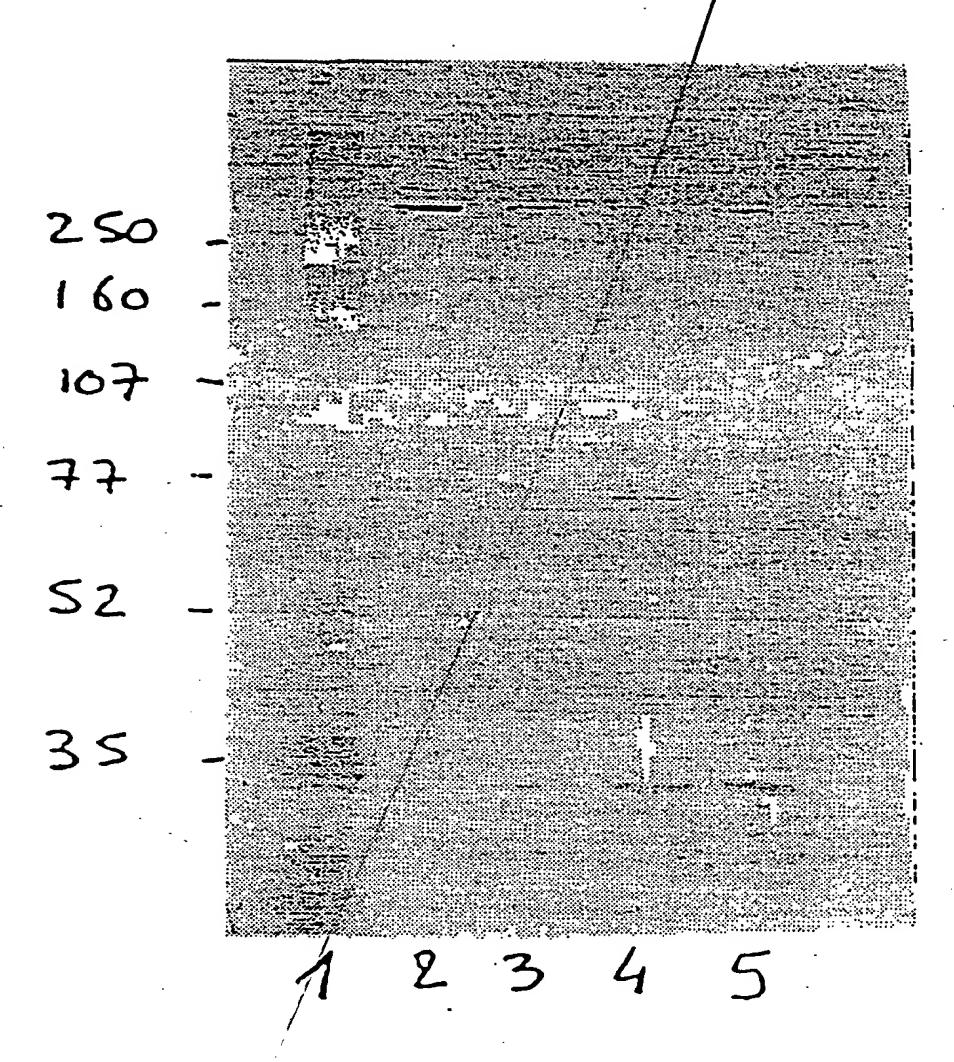


FIGURE 7
REPLACEMENT SHEET (RULE 26)

Leu Ala Ser Arg Cys Arg Cys Cys Arg Pro Leu Val Glu Ala Ala Val His Gly Trp Arg Val Glu Ala Ala Ala Gly Arg Cys Cys Arg Leu Leu Met Gly Gly Ala Cys Lys Pro Leu Pro Leu Val Glu Ala Ala Gly ** Cys Cys Cys Ala TOG TOG COT GAA COO GTO GOO GTO GTO GAG COO TOG TOG AGT COT COT TOT ACG 18 27 36 ACC AGC GCA CIT COG CAG COG CAG CAC CITC COC AGC ACC /TCA GCA GCA ACA TGC 45 5. Thr Ser Ala Leu Arg Gln Arg Gln His Leu Gly Ser Thr/Ser Ala Ala Thr Cys Pro Ala His Phe Gly Ser Gly Ser Thr Ser Ala Ala Pro Gln Gln His Ala Gln Arg Thr Ser Ala Ala Ala Ala Pro Arg Gln His Veu Ser Ser Asn Met Pro Ala Leu Leu Ile Ser Ser Ala Ser Gly Leu Gly Met /Phe Pro Pro His Glu Ser Leu Leu Phe Phe Pro Leu Leu Pro Gly Trp Gly Trp Leu Leu His Thr Asn Val Trp Cys Ser Ser His Phe Phe Arg Val Gly Val Gly Tyr Phe Thr Pro Thr *** GGT CGT TCT TCT TAC CTT CTT CGC CTG GGG TTG GGG TAT TTT CCA CCC ACA AGT 72 99 CCA GCA AGA AGA ATG GAA GAA GCG GAC CCC AAC CCC /ATA AAA GGT GGG TGT TCA Pro Ala Arg Arg Met Glu Glu Ala Asp Pro Asn Pro Ile Lys Gly Gly Cys Ser Gln Gln Glu Glu Trp Lys Lys Arg Thr Pro Thr Pro ... Lys Val Gly Val His Ser Lys Lys Asn Gly Arg Ser Gly Pro Gln Pro His Lys Arg Trp Val Phe Thr Gln Ile Ile Arg Gly Phe Val Leu Ala Leu Phe Tyr Pro Ile Lys Trp Tyr Gly Arg Phe Leu Gly Glu Ser Ser Ser Arg Leu Phe/Ile Arg Ser Arg Gly Ile Asp Glu Ser Tyr Asp Lys Arg Leu Arg Ala Cys Ser/Phe Val Pro Asp Glu Leu Ile GAG ACT TAT TAG GAA GOO TTO TGC TCG CGT TCT TTT ATG CCC TAG AAG GTT ATA 117 126 135 144 153 CTC TGA ATA ATC CTT CCG AAG ACG AGC GCA AGA AAA TAC GCG ATC TTC CAA TAT 162 Leu *** Ile Ile Leu Pro Lys Thr Ser Ala Arg Lys Tyr Gly Ile Phe Gin Tyr Ser Glu *** Ser Phe Arg Arg Arg Ala Glń Glu Asn Thr Gly Ser Ser Asn Ile Leu Asn Asn Pro Ser Glu Asp Glu Arg Lys Lys Ile Arg Asp Leu Pro Ile Ser *** Lys Ile Ile Lys Asn Asn Ala Leu/Leu Thr Ile Leu Phe Ser Ser Cys Arg Arg Asn Ser *** Lys Ile Thr Pro Ser/Ser Pro Leu Ser Ser Pro Arg Val Gly Gly Ile Gln Asn Asn *** Gln Gln Arg /Pro Pro Tyr His Pro Leu Val Phe Val CGG ATA AAC TAA TAA AAT AAC AAC CGG TCC TCC CAT TAC TCC TTC CTG CTT GTG 171 180 189 198 207 216 CCC TAT TIG ATT ATT TTA TIG TIG GCG AGG AGG GTA ATG AGG AAG GAC GAA CAC Pro Tyr Leu Ile Ile Leu Leu Leu/Ala Arg Arg Val Met Arg Lys Asp Glu His Pro Ile *** Leu Phe Tyr Cys Trp Arg Gly Gly *** *** Gly Arg Thr Asn Thr Leu Phe Asp Tyr Phe Ile Val Cly Glu Glu Glu Glu Glu Glu Glu Gly Arg Thr Pro Val Glu Leu Pro Glu Ser Ile Lys His Leu Leu Leu Ser Lys Ile Phe His Leu *** Arg Trp Pro Asn Ala Leu Lys Thr Phe Phe Cys Val Lys Leu Leu Thr Phe Glu Gly Gly Pro Thr Arg ***/Asn Gln Ser Ser Ala Ser Lys *** Tyr Leu Ser GAG TGG AGG TCC CCA AGC GAT TAA AAC ACT TCT TCG TCT GAA AAT TAT TTC ACT 234 243 252 261 270 CTC ACC TCC AGG GGT TCG CTA ATT TTG TGA AGA AGC AGA CTT TTA ATA AAG TGA Leu Thr Ser Arg Gly Ser/Leu Ile Leu *** Arg Ser Arg Leu Leu Ile Lys ***

FIGURE 8

Ser Pro Pro Gly Val Arg *** Phe Cys Glu Glu Ala Asp Phe *** *** Ser Glu

His Leu Gln Gly Phe/Ala Asn Phe Val Lys Lys Gln Thr Phe Asn Lys Val Lys

Pro Ile Gln Thr Gly Ala Ala Val Asp Leu Phe Arg Phe Ser Cys Ile Leu Leu His Tyr Lys Pro Ala Arg Gln Trp Met Ser Phe Ala Phe Pro Val Ser *** Cys Thr Thr Asn Pro His Gly Ser Gly Cys Arg Ser Leu Ser Leu Phe Leu Asp Ala TCA CCA TAA ACC CAC GCG CCA CCG TCT ACC TCT TTC CCT TTC CTT GTC TAG TCG 279 288 297 306 315 324 ACT GCT ATT TCG GTG CCC GCT GCC ACA TCG AGA AAG CGA AAG GAA CAG ATC ACC Ser Gly Ile Trp Val Pro Ala Ala Thr Ser Arg Lys Arg Lys Glu Gln Ile Ser Val Val Phe Gly Cys Pro Leu Pro His Arg Glu Ser Glu Arg Asn Arg Ser Ala Trp Tyr Leu Gly Ala Arg Cys His Ile Glu Lys Ala Lys Gly Thr Asp Gln Gln

Ile Phe Phe Val Ala Thr Phe Phe Ala Val ... Gin His Leu Thr Ser Ser Arg Phe Leu Ser Tyr Gin Leu Leu Ser Pro Leu Lys Ser Ile Ser His Pro Ala Gly Ser Tyr Leu Ile Ser Cys Tyr Leu Leu Cys Ser Vai Ser Pro Thr His Leu Glu TCT TAT TTC TTA TGA CGT CAT TTC TTC CGT TGA ATG ACT ACC TCA CAC CTC GAG 333 342 351 360 369 378 AGA ATA AAG AAT ACT GCA GTA AAG AAG CCA ACT TAC TGA TGG AGT GTG GAG CTC Arg Ile Lys Asn Thr Ala Val Lys Lys Ala Thr Tyr ... Trp Ser Val Glu Leu Glu ... Arg Ile Leu Gin ... Arg Arg Gln Leu Thr Asp Gly Val Trp Ser Ser Asn Lys Glu Tyr Cys Ser Lys Glu Gly Asn Leu Leu Met Glu Cys Gly Ala Pro

Ser Arg Leu Ser Leu Pro Thr Val Gln Arg Ser Ser His Thr Gly Gln Gln Leu Leu Asp *** Pro Cys Arg Leu Ser Arg Asp/Val Ala Thr Leu Val Lys Asn Ser Ile Glu Pro Val Val Ser His Gly Thr *** Gln Gln Ser Tyr Arg Thr Pro GAT CTA GAG TCC CTG TTG CCT CAC TCG ACA GAT GAC GAC ACT CAT GGA ACA ACC 387 396 405 414 423 432 CTA GAT CTC ACG GAC AAC GGA GTG ACC TGT CTA CTG CTG TGA GTA CCT TGT TCG Leu Asp Leu Arg Asp Asn Gly Val Thr Cys Leu Leu Leu *** Val Pro Cys Trp Ile Ser Gly Thr Thr Glu *** Pro Val Tyr Cys Cys Glu Tyr Leu Val Gly Arg Ser Gln Gly Gln Arg Ser Asp Leu Ser Thr Ala Val Ser Thr Leu Leu Glu

Ala Pro Thr Gln His Gly Asn Cys Leu Leu Val Arg Tyr Arg Lys Asp Ser Ile Leu Pro Leu Arg Thr Val Thr Ala Ser Cys Cys Gly Thr Val Asn Thr Leu Phe Ser Arg Ser Asp Pro Ser Arg Gln Leu Ala Ala Gly Gln Leu Thr Gln ... Phe TCT CCC CCT CAG ACC ACT CCC AAC GTC TCG TCG TCG GAC ATT GCA AAC AGT CTT 441 450 459 468 477 486 AGA CCG GGA GTC TCG TGA CCG TTG CAG AGC ACC CTG TAA CGT TTG TCA GAA ARG AGG GGA GTC TCG TGA CCG TTG CAG AGC ACC CTG TAA CGT TTG TCA GAA ARG ALG GLy Val Trp ... Pro Leu Gln Ser Ser Thr Leu ... Arg Leu Ser Glu Glu Arg Glu Ser Gly Asp Arg Cys Arg Ala Ala Pro Cys Asn Val Cys Gln Lys Ser Gly Ser Leu Val Thr Val Ala Glu Gln His Pro Val Thr Phe Val Arg Asn

Glu Ala Pro Gln Ser Phe Lys Gln Phe His Ala Pro Phe His Leu Leu Thr Ile Lys Arg Pro Ser Ala Ser Ser Lys Phe Thr Leu Pro Phe Ile Cys Phe Arg Ser Asn Gly Arg Ala Pro Gln Val Lys Ser Leu Ser Arg Ser Phe Ala Ser Ala His TAA AGG CGC CCG ACC GAG TTG AAA ACT TTC ACT CCC CCT TTT ACG TCT TCG CAC 495 504 513 522 531 540 ATT TCC CCG CCC TCG CTG AAC TTT TGA AAG TGA CCG GGA AAA TCC AGA ACC GTG Lie Ser Ala Gly Trp Leu Asn Phe *** Lys *** Ala Gly Lys Cys Arg Ser Val Phe Pro Arg Ala Gly *** Thr Phe Glu Ser Glu Arg Glu Asn Ala Glu Ala *** Phe Arg Gly Leu Ala Glu Leu Leu Lys Val Ser Gly Lys Met Gln Lys Arg Asp

FIGURE 8 (continuation 1)

Pro Leu Ser Ile Tyr Val Asp Asn His Pro Trp Arg Pro Thr Thr Phe Ala Phe Gln Phe Val Leu Thr Cys Thr Met Thr Pro Gly Gly Pro His Pro Leu Leu Leu Asn Ser Ser "His Val Arg "Gln Pro Ala Val Gln Thr His Tyr Phe Cys

TAA CCT TCT GAT TAC ATG TCC ACT AAC ACC CCG GTG GAC CCA CAC CAT TTT CGT 549 558 567 576 585 594

ATT GGA AGA CTA ATG TAC ACG TCA TTG TCG GCC CAC GTG GCT GTG GTA AAA GCA

Ile Gly Arg Leu Met Tyr Thr Ser Leu Trp Gly His Leu Gly Val Val Lys Ala Leu Glu Asp "Cys Thr Arg His Cys Gly Ala Thr Trp Val Trp "Lys Gln Trp Lys Thr Asn Val His Val Ile Val Gly Pro Bro Gly Cys Gly Lys Ser Lys

Pro Ser Ser Ile Lys Cys Val Arg Phe Gly Cys Val Pro Phe Trp Arg Ser Val His Ala Ala Leu Lys Ala Ser Gly Ser Val Val Jyr Gln Phe Gly Gly Leu Phe Ile Pro Gln *** Asn Gln Leu Gly Pro Phe Trp Met Ser Ser Val Val *** Phe TTA CCC GAC GAT TAA AAC GTC TCC GCC TTT GCT GTA TGA CCT TTG GTG GAT CTT 603 612 621 630 639 648 AAT GCG CTG CTA ATT TTG CAG ACC CCG AAA CCA CAT ACT GGA AAC CAC CTA GAA Asn Gly Leu Leu Ile Leu Gln Thr Arg Lys Pro His Thr Gly Asn His Leu Glu Met Gly Cys *** Phe Cys Arg Pro Gly Asn/His Ile Leu Glu Thr Thr *** Lys Trp Ala Ala Asn Phe Ala Asp Pro Glu Thr Thr Tyr Trp Lys Pro Pro Arg Asn

Leu Pro Pro Ile Thr Val Met Thr Phe Phe His Asn Asn Asn Ile Val Lys Ile
Leu His His Ser Pro *** Trp Pro Ser Ser Thr Thr Thr Ile Ser Ser Lys

Cys Thr Thr Pro His Asn Gly His His Leu Leu Pro Gln *** Gln His Ser Lys

TGT TCA CCA CCC TAC CAA TCG TAC CAC TTC TTC ACC AAC AAT AAC TAC TGA AAA
657 666 675 684 693 702

ACA AGT GGT GGG ATG GTT ACC ATG GTG AAG AAG TCG TTG TTA TTG ATG ACT TTT

Thr Ser Gly Gly Met Val Thr Met Val Lys Lys Trp Leu Leu Leu Met Thr Phe
Gln Val Val Gly Trp Leu Pro Trp *** Arg Ser Gly Cys Tyr *** Leu Leu
Lys Trp Trp Asp Gly Tyr His Gly Glu Glu Val Val Val Ile Asp Asp Phe Tyr

Ala Pro Gln Gly Pro Ile Ile ... Gln Ser Gln Thr Ile Ser Ile Trp Gln Ser Pro Gln Ser Gly Gln Ser Ser Arg Ser Leu Ser His Ser Arg Tyr Gly Asn Val His Ser Ala Ala Arg Pro His Asp Val Ser Val Thr His Asp Ile Asp Met Ser TAC CGA CCG ACG GGA CCC TAG TAG ATG ACT CTG ACA CAC TAG CTA TAG GTA ACT 711 720 729 738 747 756

ATG GCT GGC TGC CCT GGG ATG ATC TAC TGA GAC TGT GTG ATC GAT ATC CAT TGA

Met Ala Gly Cys Pro Gly Met Ile Tyr ... Asp Cys Val Ile Asp Ile His ...

Trp Leu Ala Ala Leu Gly ... Ser Thr Glu Thr Val ... Ser Ile Ser Ile Asp Gly Trp Leu Pro Trp Asp Asp Leu Leu Arg Leu Cys Asp Arg Tyr Pro Leu Thr

Tyr Leu Ser Phe Thr Ser Ser Tyr Arg Lys Gln Gly Ala Thr Asn Gln Asn Gly Thr Ser Val Leu Pro Pro Val Thr Gly Lys Lys Ala Arg Leu Ile Arg Ile Val Gln Leu Ser ** Leu His Phe Gln Val Lys Lys Pro Gly Cys Tyr Glu Ser ** GAC ATC TCT GAT TTC CAC CTT GAC ATG GAA AAA ACC GGG CGT CAT AAG ACT AAT 765 / 774 783 792 801 810 CTG TAG AGA CTA AAG GTG GAA CTG TAC CTT TTT TGG CCC GCA GTA TTC TGA TTA Leu *** Arg Leu Lys Val Glu Leu Tyr Leu Phe Trp Pro Ala Val Phe *** Leu Cys Arg Asp *** Arg Trp Asn Cys Thr Phe Phe Gly Pro Gln Tyr Ser Asp Tyr Val Glu Thr Lys Gly Gly Thr Val Pro Phe Leu Ala Arg Ser Ile Leu Ile Thr

FIGURE 8 (continuation 2)

Ala Ile Leu Gly Arg Gln Phe Pro Val Gly ** Ser Ser Asp Trp Ser Tyr Phe Leu Leu ** Val Gly Asn Ser His Tyr Glu Glu Val Ala Thr Gly Ala Thr Ser Trp Cys Asp Ser Gly Thr Pro Ile Thr Ser Arg Leu Gln Gln Gly Leu Gln Leu GT CGT TAG TCT GGG GCA ACC TTA CCA TGA GCA GTT GAC GAC ACC GTC GAC ATC 819 828 837 846 855 864 CCA GCA ATC AGA CCC CGT TGG AAT GGT ACT CCT CAA CTG CTG TCC CAG CTG TAG Pro Ala Ile Arg Pro Arg Trp Asn Gly Thr Pro Gln Leu Leu Ser Gln Leu Gln Gln Ser Asp Pro Val Gly Met Val Leu Leu Asn Cys Cys Pro Ser Cys Arg Ser Asn Gln Thr Pro Leu Glu Trp Tyr Ser Ser Thr Ala Val Pro Ala Val Glu

Ser Lys Ile Pro Pro Asn Ser Gly Gln Tyr Lys Pro Leu Ile Ser Cys Phe Leu Ala Arg "Arg Leu Ile Val Glu Lys Thr Asn Gln Phe Phe Ala Val Ser Cys Leu Glu Lys Asp Ser Ser "Lys Arg Pro Ile Lys Ser Ser His "Leu Val TTC GAG AAA TAG CCT CCT AAT GAA GGA ACC ATA AAA CCT TCT TAC GAT GTC TTG 873 882 891 900 909 918

AAG CTC TTT ATC GGA GGA TTA CTT CCT TGG TAT TTT GGA AGA ATG CTA CAG AAC

Lys Leu Phe Ile Gly Gly Leu Leu Pro Trp Tyr Phe Gly Arg Met Leu Gln Asn Ser Ser Leu Ser Glu Asp Tyr Phe Leu Gly Ile Leu Glu Glu Cys Tyr Arg Thr Ala Leu Tyr Arg Arg Ile Thr Ser Leu Val Phe Trp Lys Asn Ala Thr Glu Gln

Gly Arg Leu Phe Pro Ala Leu Glu Asp/Gly Lys Gly Gly Trp Ala Arg Phe Lys Asp Val Ser Ser Pro Pro Trp Asn Thr/Val Arg Glu Gly Gly His Gly Ser Asn Ile Trp Pro Pro Leu Pro Gly Thr Arg ... Gly Lys Gly Gly Met Gly Gln Ile TTA CCT CCC TTC CCC CCG TCA ACC ACT CCC AAA CCC CCC CTA CCC CTC ATTA 927 936 945 954 963 972 AAT CCA CCC ACG AAG CCC ACT TCC TCC TCC CCC CAT CCC CTG AAT ASn Pro Arg Arg Lys Gly Ala Ser/Ser Ser Pro Phe Pro Pro His Ala Leu Asn Ile His Gly Gly Arg Gly Pro Val Arg His Pro Phe Pro Pro Mct Pro ... Ile Ser Thr Glu Glu Gly Gly Gln/Phe Val Thr Leu Ser Pro Pro Cys Pro Glu Phe

Trp Ile Phe Tyr Ile Val Ser Asp Lys Lys Asp Ser Arg Leu Pro Lys

Gly Tyr Ser Ile Phe ... Gln Thr Lys Lys Ile Val Glu Tyr His Asn Lys Asn

Glu Met His Phe Leu Asn Ser Leu Arg Lys ... Lys Thr Ile Thr Lys Ile

AAG GTA TAC TTT ATT TAA TGA CTC AGA AAA AAT AGT GAA GCA TTA CCA AAA ATA

981 990 999 1008 1017 1026

TTC CAT ATG AAA TAA ATT ACT GAG TCT TTT TTA TCA CTT CGT AAT GGT TTT TAT

Phe His Met Lys ... Ile Thr Glu Ser Phe Leu Ser Leu Arg Asn Gly Phe Tyr

Ser Ile ... Asn Lys Leu Leu Ser Leu Phe Tyr His Phe Val Met Val Phe Ile

Pro Tyr Glu Ile Asn Tyr ... Val Phe Phe Ile Thr Ser ... Trp Phe Leu Leu

Glu Asn Leu Thr Leu His Pro Thr Lys Leu Ile Leu Asn Glu Ser Asn Tyr Met Asn Met Leu Pro / Thr Pro Pro Arg Phe Ile Arg Gln Ile Thr Cys Ile Thr Pro Pro Asn Leu Pro Pro Asp Lys Phe Asn Phe Glu Arg Phe Gln Val ATA AGT AAT TCE CAA TTC ACC CCC CAG AAA TTT TAA TTT AAG AGA CTT AAC ATG 1035 1044 1053 1062 1071 1080

TAT TCA TTA AGG GTT AAG TGG GGG GTC TTT AAA ATT AAA TTC TCT GAA TTG TAC Tyr Ser Leu Arg Val Lys Trp Gly Val Phe Lys Ile Lys Phe Ser Glu Leu Tyr Ile His Gly Leu Ser Gly Gly Ser Leu Lys Leu Asn Ser Leu Asn Cys Thr Phe Ile Lys Gly Val Gly Gly Leu Asn VIle Leu Ile Val His

FIGURE 8 (continuation 3)

Cys Pro *** Val Ser Ile Thr Asn Arg Thr Thr Tyr Val Thr Lys Ser Arg Leu Val His Asn Cys Pro Tyr Gln Ile Gly Pro Arg Ile Tyr Gln Lys Arg Val Cys Tyr Met Thr Val Arg Ile Asn Tyr Glu Gln Asp Tyr Ile Ser Asn Glu Phe Ala TAT GTA CCA ATG TCC CTA TAA CAT AAG GAC CAG CAT ATA TGA CAA AAG CTT CCG 1089 1098 1107 1116 1125 1134 ATA CAT GGT TAC ACG GAT ATT GTA TTC CTG GTC GTA TAT AGT GTT TTC GAA CCC Ile His Gly Tyr Thr Asp Ile Val Phe Leu Val Val Tyr Thr Val Phe Glu Arg Tyr Met Val Thr Arg Ile Leu Tyr Ser Trp Ser Tyr Ile/Leu Phe Ser Asn Ala Thr Trp Leu His Gly Tyr Cys Ile Pro Gly Arg Ile Tyr Cys Phe Arg Thr Gln

Ala Ser Ala *** Thr Thr *** Met Glu Leu Leu Lys Tyr Asp *** Gly Cys Ser His Arg Pro Arg Arg Pro Arg Cys Lys Trp Cys Asn Thr Thr Glu Ala Val Ala Thr Gly Leu Gly Val His Asp Val Asn Gly Ala Thr Gln Leu Arg Leu Trp Leu TCA CCC CTC CCC ATG CAC CAG ATG TAA ACG TCG TCA AAC ATC AGA GTC CCT GTC 1143 1152 1161 1170 1179 1188

AGT CCC GAG GCC TAC GTC GTC TAC ATT TCC AGC AGT TTC TAG TCT CAG CCA CAG

Ser Ala Glu Ala Tyr Val Val Tyr Ile Ser Ser Ser Leu ** Ser Gln Pro Gln Val Pro Arg Pro Thr Trp Ser Thr Phe Pro Ala Val Cys Ser Leu Ser His Ser Cys Arg Gly Leu Arg Gly Leu His Phe Gln Gln Phe Val Val Ser Ala Thr Ala

Thr Glu Lys Thr Thr Gln Asn Ser Thr Ile Leu Leu Ser Ile Ser Leu Asn Pro Lys Lys Gln Gln Lys Thr Pro Leu Leu Tyr His Phe Arg Pro Cys Thr Gln Asn Arg Lys Asn Asn Pro Gln Phe Tyr Asp Ile Thr Phe Asp Leu Val Pro GAC CAA AGA AAA CAA ACC AAC CTT CAT TAG TTA TCA CTT TAG ATC CTG TCC 1197 1206 1215 / 1224 1233 1242 CTG GTT TCT TTT GTT GTT TGG TTG GAA GTA ATC AAT AGT GAA ATC TAG GAC ACG Leu Val Ser Phe Val Val Trp Leu Glu Val Ile Asn Ser Glu Ile Asp Arg Trp Phe Leu Leu Leu Phe Gly Trp Lys Ser Ile Val Lys Ser Arg Thr Gly Gly Phe Phe Cys Cys Leu Val Gly Ser Asn Gln Asn Leu Gly Gln Val

Pro Pro Leu Thr Gly Pro Thr Thr Pro Ser Pro Ser Pro Pro Ile Ala Pro Gln Pro Tyr Leu Val Pro Leu Pro Leu Leu Leu Ala Pro Asn His Tyr Pro Pro Lys Pro Thr Phe Tyr Arg Ser His Tyr Ser Phe Pro Gln Thr Ile Thr His Arg AAA CCC CCA TIT CAT CCC CCT CAC CAT CCT CTT CCC GAC CCA ATA CCA TAC CCC 1251 1260 1269 1278 1287 1296

TIT CCC CCT AAA GTA CCC CCA GTC GTA CCA GAA CCC CTC CCT TAT CCT ATC CCC Phe Gly Gly Lys Val Pro Gly Val Val Gly Glu Gly Leu Gly Tyr Gly Met Ala Leu Gly Val Lys Tyr Arg Glu Trp Gly Arg Arg Arg Arg Ala Gly Leu Trp Tyr Gly Gly

Pro Thr Thr *** Met Pro Thr Met Pro Ser Pro Gln Pro Arg Gln *** Leu Thr Leu Leu Leu Lys Cys Leu Pro *** Leu His Pro Ser His Gly Lys Asn Cys Leu Ser Ser Tyr Asn Val Tyr Pro Asp Tyr Thr Leu Ala Thr Ala Lys Thr Val Phe CCT CCT CAT CAA ATG TAT CCC CAG TAT CCA CTC CCG ACA CCG GAA ACA ATG TTT 1305 1314 1323 1332 1341 1350 CGA GGA GTA GTT TAC ATA GGG GTC ATA GGT GAG CGC TGT CCC CTT TGT TAC AAA Gly Gly Val Val Tyr Ile Gly Val Ile Gly Glu Gly Cys Gly Leu Cys Tyr Lys Glu Glu *** Phe Thr *** Gly Ser *** Val Arg Ala Val Ala Phe Val Thr Lys Arg Ser Ser Leu His Arg Gly His Arg *** Gly Leu Trp Pro Leu Leu Gln Ser

FIGURE 8 (continuation 4)

Ile Met *** Phe Leu Leu Val Pro Ala Trp Glu Gly Thr Val Arg Pro Ser Arg

Arg Phe Tyr Cys Cys Gln Leu Gly Ser Gly Gln ** Gly Pro His Asp

Asn Asp Asp Leu Ile Val Ala Ser Ser Gly Val Gly Arg Asp Gly Gln Thr Ile

CAA TAG TAG ATT TTA TTG TCG TCA CCT CCC GTG ACG CGA CAG TCG GAC CCA CTA

1359

1368

1377

1386

GTT ATC ATC TAA AAT AAC ACC ACT GGA CCC CAC TCC CCT GTC ACC CTG GGT GAT

Val Ile Ile ** Asn Asn Ser Thr Gly Ala His Ser Pro Val Thr Leu Gly Asp

Leu Ser Ser Lys Ile Thr Ala Leu Glu Pro Thr Pro Leu Ser Pro Trp Val Ile

Tyr His Leu Lys ** Gln His Trp Ser Pro Leu Pro Cys His Pro Gly ** Ser

Pro Ala Pro Gly Ser Asn Leu Arg Leu Arg Glu Glu Thr Thr Asn Leu Pro Pro Leu Leu Ala Leu Ile Gly Gly Lys Lys Asn Gln Leu Ile Leu Pro Ser Cys Pro Trp Phe Glu Val Lys Val Lys Arg Ile Arg Tyr Tyr Glu Phe CCC CCT CGT CCC GGT CTT AAG TTG GAA TTG GAA AGA ATA AGA CAT CAT AAG TTT 1413 1422 1431 1440 1449 1458 CCG GGA GCA GCG CCA GAA TTC AAC CTT AAC CTT TCT TAT TCT GTA GTA TTC AAA Arg Gly Ala Gly Pro Glu Phe Asn Leu Asn Leu Ser Tyr Ser Val Val Phe Lys Gly Glu Gln Gly Gln Asn Ser Thr Leu Thr Phe Leu Ile Leu Tyr Ser Lys Gly Ser Arg Ala Arg Ile Gln Pro Pro Phe Leu Phe Cys Ser Ile Gln Arg

Cys Leu Ala Pro Thr Gln Gly Gly Glu Gln Pro Phe Phe Thr Met Leu Ile Ser Ala Cys Leu Pro Pro Lys Val Gly Arg Arg Pro Ser Ser Leu Tyr Gln Pro Val Ser Arg Pro Asn Ser Gly Gly Gly Pro Pro Leu Phe Asp Asn Ile Asn CCC GTG TCT CCC CCC CAA ACT GCG GCG ACG ACC CCC TTC TTT CAG TAA TTA TAA 1467 1476 1485 1494 1503 1512 CCC CAC AGA GCG GGG GTT TGA CCC CCC TCC TCG GCG AAG AAA GTC ATT AAT ATT Gly His Arg Ala Gly Val Pro Pro Ser Trp Gly Lys Lys Val Ile Asn Ile Gly Thr Glu Arg Gly Phe Asp Pro Pro Pro Gly Gly Arg Lys Ser Leu Ile Leu Ala Gln Ser Gly Gly Leu Thr Pro Leu Leu Gly Glu Glu Ser His Tyr

Asp *** Thr Trp Arg Gly Pro Pro Arg Glu Ser Gln Pro Glu Ser Ser Leu Ile Glu Asp His Gly Gly Leu Leu Ala Asn Gln Ser His Asn Ala Gln Cys Phe Arg Met Met Asp Val Ala Trp Ser Pro Thr Arg Val Thr Thr Arg Lys Val CTT AGA GTA GTA CAG GTG GCG GCT CCT CCC GCA AGA CTG ACA CCA AGC GAA CTG 1521 1530 1539 1548 1557 1566 GAA TCT CAT CAT GTC CAC CGC CCA GGA GCG CGT TCT GAC TGT GCT TCG CTT GAC Glu Ser His His Val His Arg Pro Gly Gly Arg Ser Asp Cys Gly Ser Leu Asp Asn Leu Ile Met Ser Thr Ala Gln Glu Gly Val Leu Thr Val Val Arg Leu Thr Ile Ser Ser Cys Pro Pro Pro Arg Arg Ala Phe *** Leu Trp Phe Ala *** Gln

Ile Asp Ser Pro Ala Pro Ser Ala Pro Thr Ser Ser Ala Met Lys Gly Glu Gly Tyr Ile Arg Leu His Pro Leu Pro Pro His Gln Leu His Trp Lys Glu Lys Glu Thr Tyr Gly Phe Thr Arg Ser Leu Arg Thr Asn Phe Ile Gly Asn Lys Arg Arg TCA TAT AGG CTT CCA CCC CCT CTC CCC CCA CAA CTT CTA CCC TAA AAA CGA AGA 1575 1584 1593 1602 1611 1620 AGT ATA TCC GAA GGT GCG GGA GAG GCG GGT GTT GAA GAT GCC ATT TTT CCT TCT Ser Ile Ser Glu Gly Ala Gly Glu Ala Gly Val Glu Asp Ala Ile Phe Pro Ser Val Tyr Pro Lys Val Arg Glu Arg Arg Val Leu Lys Met Pro Phe Phe Leu Leu Tyr Ile Arg Arg Cys Gly Arg Gly Gly Cys *** Arg Cys His Phe Ser Phe Ser

FIGURE 8 (continuation 5)

Leu Ile Ala Ala Pro Ala Thr Asp Glu Glu Glu Thr Val Gly Gly Gln Ile Arg
Trp Ser Pro Gln Pro Pro Pro Thr Lys Lys Lys Pro Leu Ala Glu Lys Ser Val
Gly Leu His Ser Arg Pro Arg His Arg Arg Arg Arg Tyr Arg Arg Arg Pro Tyr

CGG TTC TAC CGA CGC CCC CGC CAC AGA AGA AGA AGC CAT TGC CGA GGA ACC TAT

1683

1692

1701

1710

1719

1728

CCC AAG ATG CCT GCG GGG GCG GTG TCT TCT TCT TCG GTA ACG CCT CCT TGG ATA

Ala Lys Met Ala Ala Gly Ala Val Ser Ser Ser Ser Val Thr Pro Pro Trp Ile

Pro Arg Trp Leu Arg Gly Arg Cys Leu Leu Leu Arg

Arg Leu Leu Gly Tyr

Gln Asp Gly Cys Gly Gly Gly Val Phe Phe Phe Gly Asn Ala Ser Leu Asp Thr

Asp Tyr Arg Phe Val Phe Ser Thr Arg Gln Leu Tyr
Thr Met Asp Ser Phe Ser Leu Leu Ala Ser Tyr Thr Asn

GCA GTA TAG ACT TTT GCT TTC TTC ACG CGA CAT TCA TAA 5.

1737

1746

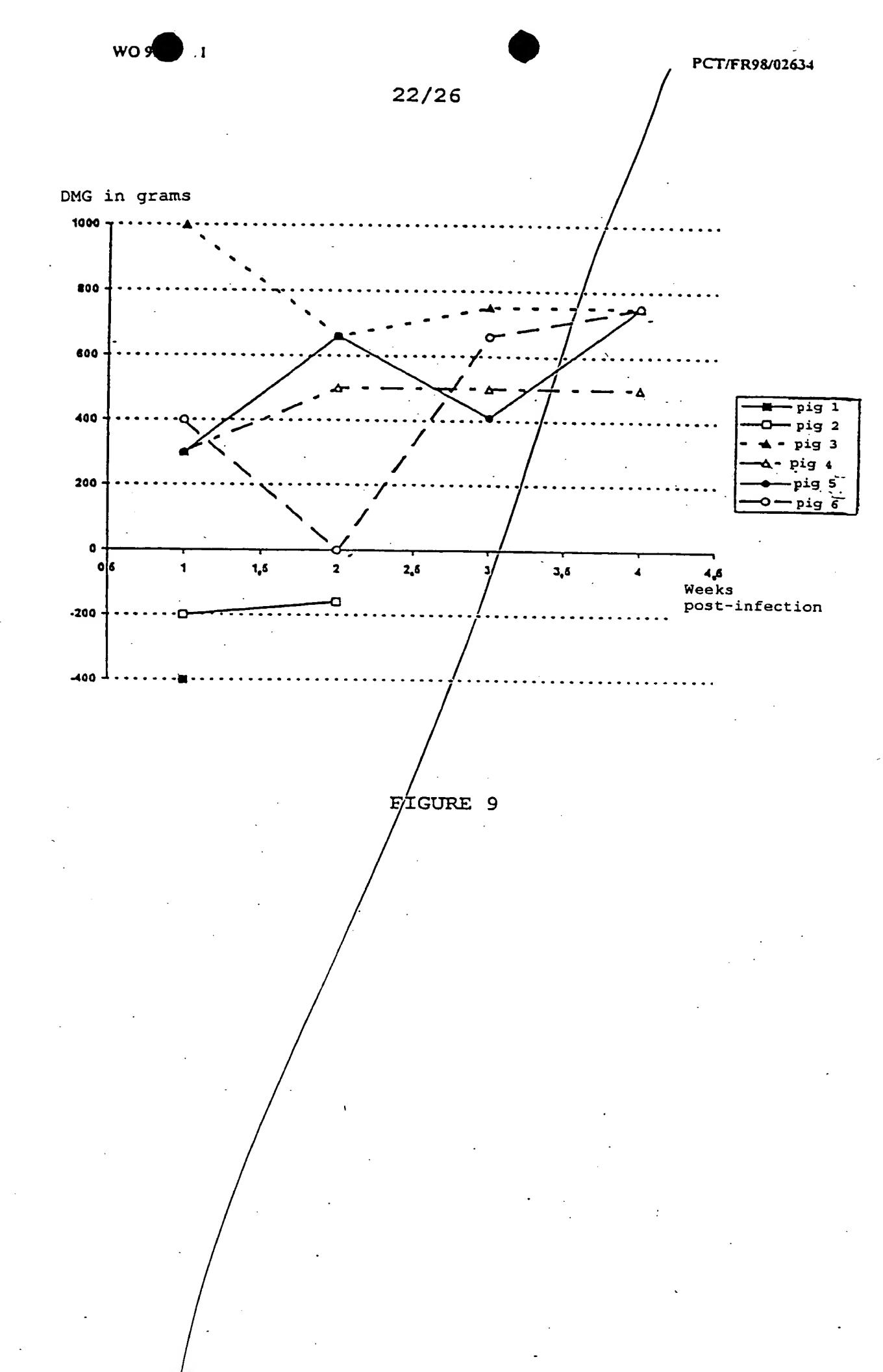
1755

1764

CGT CAT ATC TGA AAA CGA AAG AAG TGC GCT GTA AGT ATT 3.

Arg His Ile ... Lys Arg Lys Lys Cys Ala Val Ser Ile
Val Ile Ser Glu Asn Glu Arg Ser Ala Leu ... Val
Ser Tyr Leu Lys Thr Lys Glu Val Arg Cys Lys Tyr

FIGURE 8 (continuation 6)



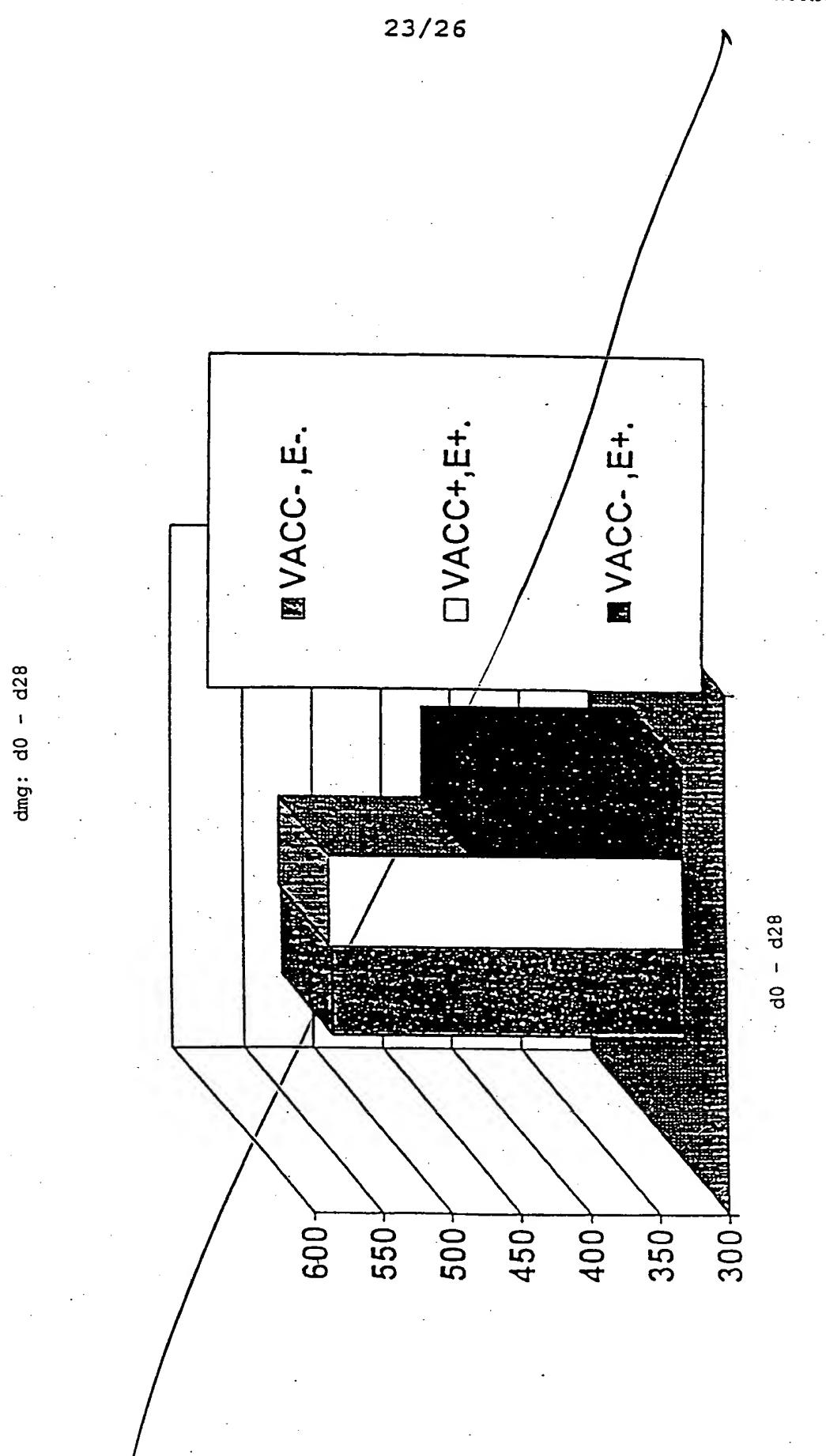


FIGURE 10

& HYPERTHERMIA >

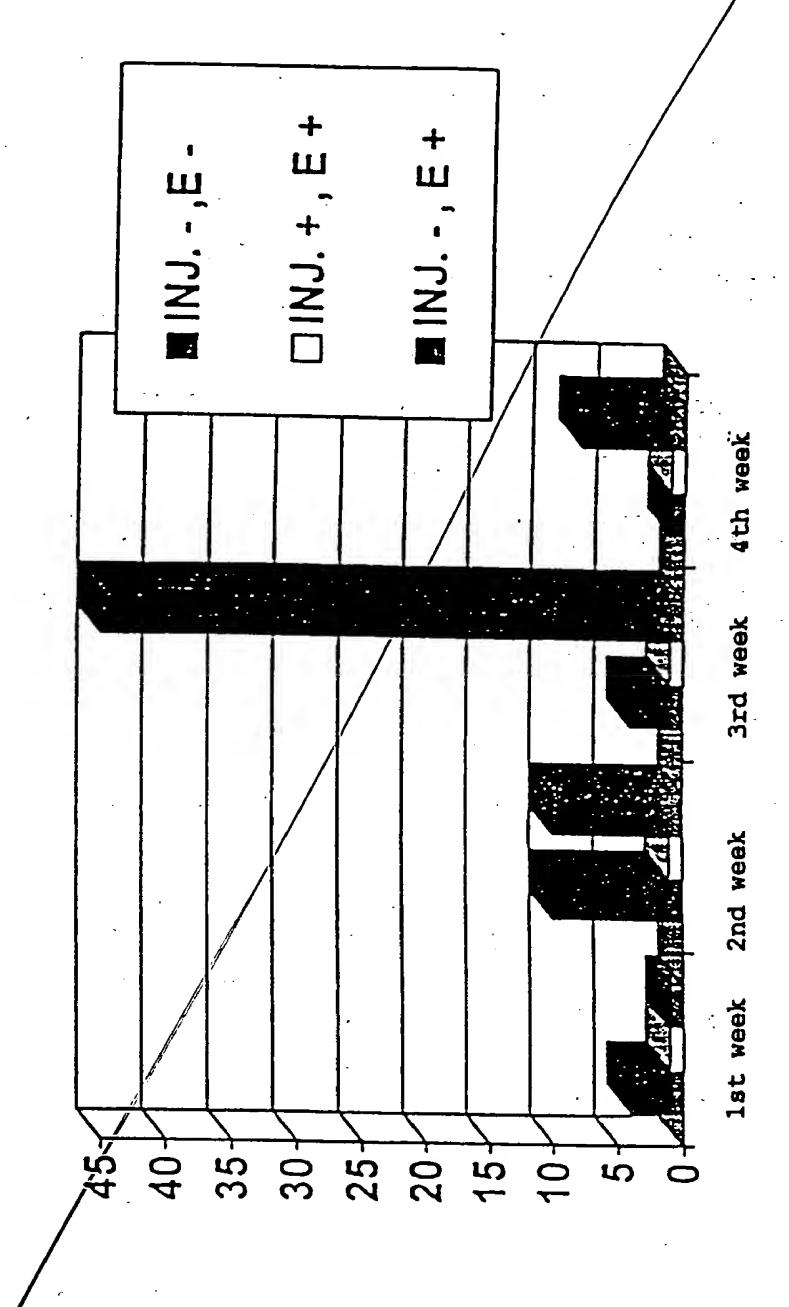
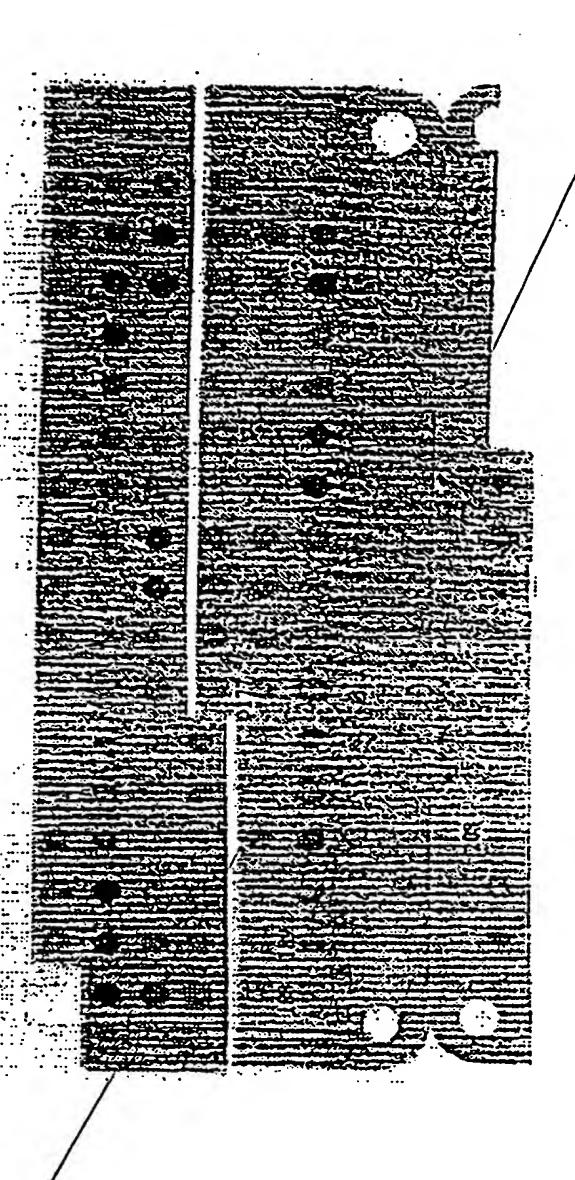


FIGURE 11





spot No. 104 to 1

v odši

GYALQNATTA

peptide 208

GTAFENSIYD

TAGNVDHVGL

THTNVEHTGL

KRNQLWLHLN

OTIBMEQPNN

RYFTPKPELD

TIROPFTYHS

pcva DPYINYSSRH

150

TITOPESYHS

DPYVNYSSRH

pcvB

RYFTPKPVLD

FTIDYFOPNN

KRNQLWLRLQ

RGGHSQPSWN RTTVRTPSWA LSREFVLTI. LSRTFGYTVK 50 RRKTGI FNSR RRKNGI ENTR HP. RHRYRW HPAFRNRYRW QILRRRPWLV NILRRRPYLV RRHRPRSHLG RRTRPRSHLG MTYPRRAYRR MTWPRRRYRR pcvA pcvB

101 100

peptides 188 to 189

FVTPSTNLAY **FVTKATALTY** GSSAVILDDN GSTVVILDAN SPITQGDRGV DPITSNORGV RKAKYEFYPR RKVKVEFWPC RSVPFEYYRI FLPPSGGRNP LPLPFQYYRI FLPPGGGSNP peptide 177 VDMMRFNIND VNELRENIGO pcvB pcvA

peptide 121

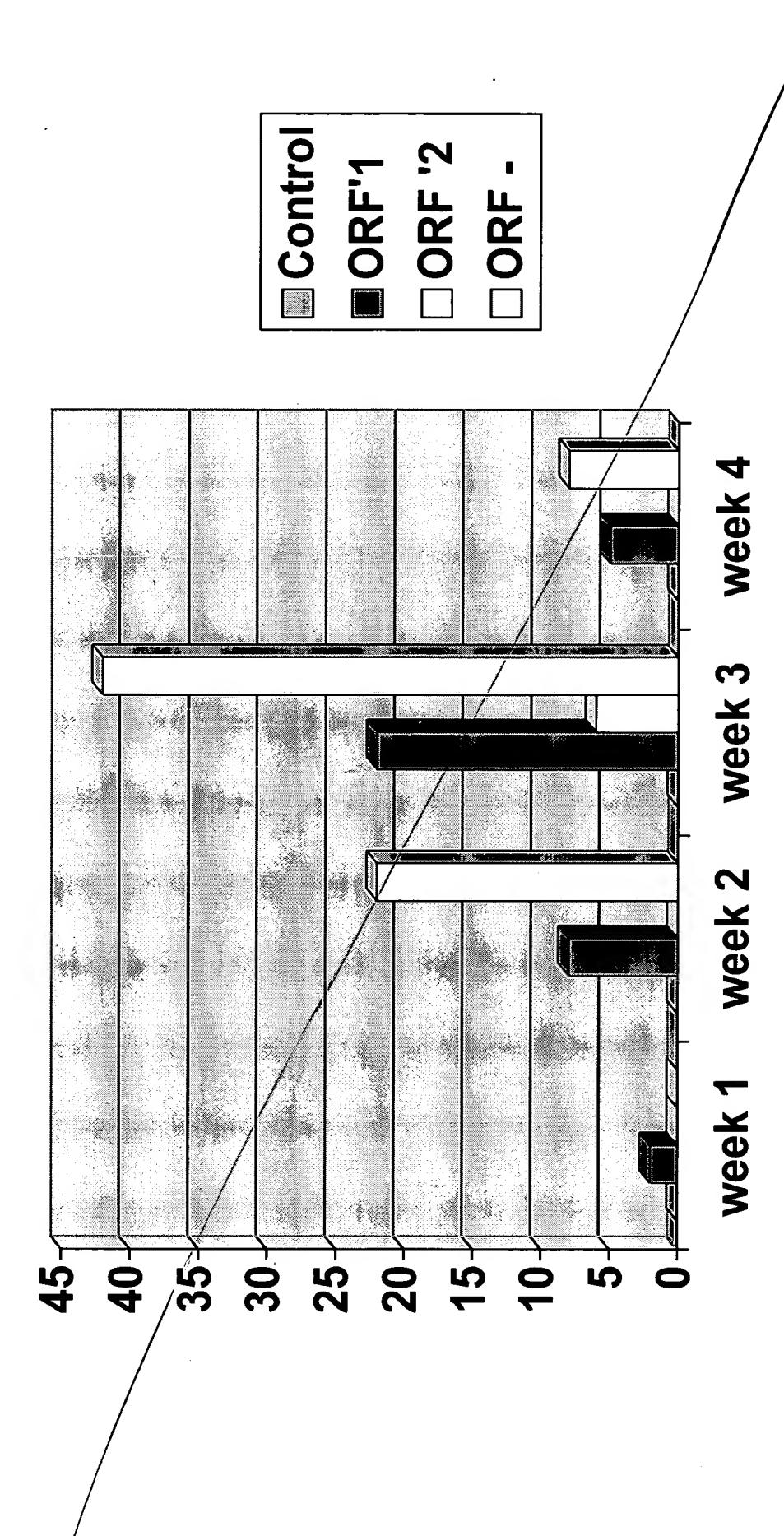
133 to 132 peptides

> PPLNP 235 P. LNE VQFREFILKD VQFREFNFKD QNYVVRLTY **QEYNIRVTMY** pcvA pcvB

peptide 152

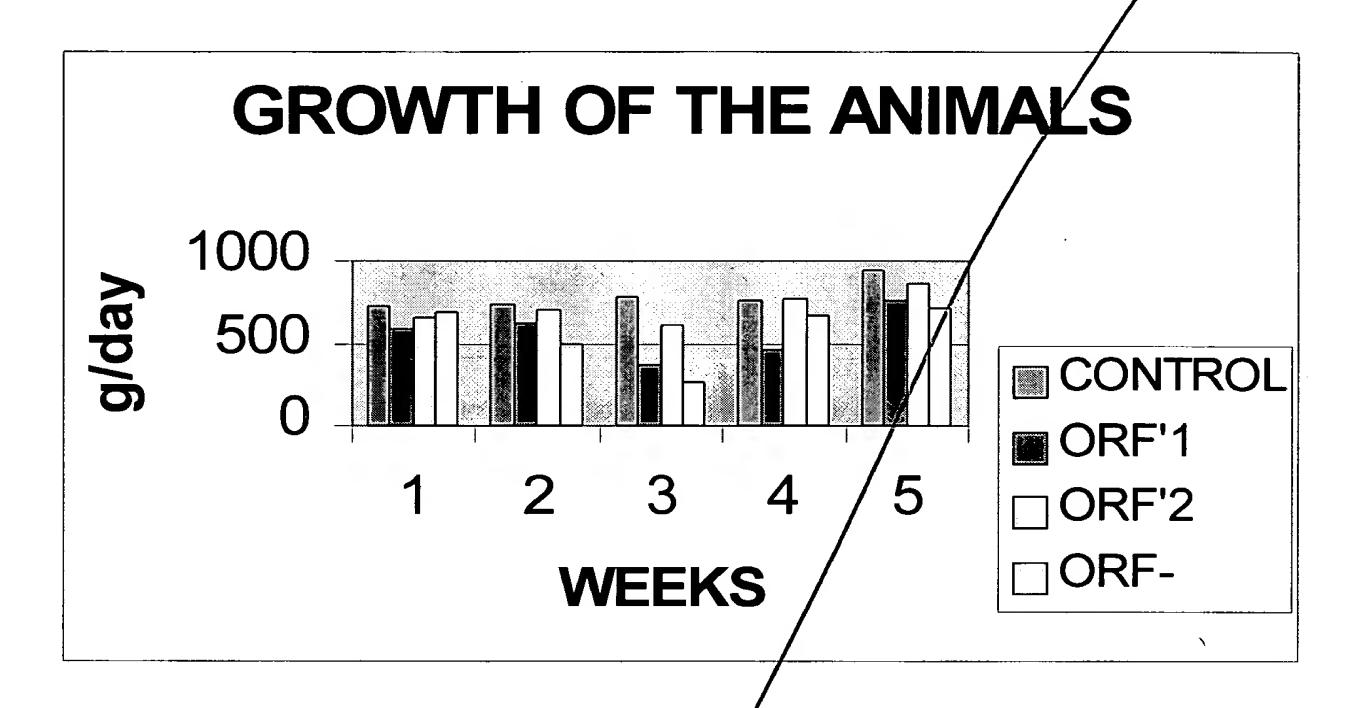
FIGURE

Figure 14



Percent hyperthermia (> 40.5°C), Control (not vaccinated and not challenged), ORF'1 (vaccinated and challenged), ORF'2 (vaccinated and challenged), ORF- (not vaccinated, challenged)

Figure 15



Control (not vaccinated and not challenged), ORF'1(vaccinated and challenged) ORF'2 (vaccinated and challenged), ORF- (not vaccinated, challenged)

Figure 16

Immunoperoxidase staining of PK15 cells at 24 h post-transfection with the pcDNA3/ORF'2 plasmid. Expression of PCV-B ORF'2 was confirmed by IPMA following incubation in the presence of the swine anti-PCV-B monospecific serum.

